

1 **Biological sample collection to advance research and treatment: a Fight Osteosarcoma Through**
 2 **European Research (FOSTER) and Euro Ewing Consortium (EEC) statement.**

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111 **Running Title:** EEC and FOSTER sample statement

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115 **STATEMENT OF TRANSLATIONAL RELEVANCE**

116 Most patients with osteosarcoma and Ewing sarcoma have minimal sampling performed at clinical
117 presentation, sufficient for diagnosis but not for comprehensive molecular analysis. Mechanistic
118 understanding of tumorigenesis, metastasis and treatment resistance has progressed little. Standard
119 management involves upfront biopsy, frequently by an interventional radiologist, followed by
120 chemotherapy +/- definitive resection, by which time post-treatment necrotic tumour may be less
121 informative for cellular analysis and model generation. Few patients have fresh or frozen tissue stored
122 for patient-specific or unspecified molecular research. Treatment has changed little in decades and
123 outcomes are poor. Here, the European osteosarcoma and Ewing sarcoma patient and professional
124 communities set out minimum standards for tissue sampling, sufficient for histological and molecular
125 evaluation and for all patients to have the opportunity to donate samples for research. The proposed
126 core samples will facilitate a revolution in biologically rational treatment of paediatric-type bone
127 sarcomas.

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140 **ABSTRACT**

141 Osteosarcoma and Ewing sarcoma are bone tumours mostly diagnosed in children, adolescents and
142 young adults. Despite multi-modal therapy, morbidity is high and survival rates remain low,
143 especially in the metastatic disease setting. Trials investigating targeted therapies and
144 immunotherapies have not been ground-breaking. Better understanding of biological subgroups, the
145 role of the tumour immune microenvironment, factors that promote metastasis and clinical biomarkers
146 of prognosis and drug response are required to make progress. A prerequisite to achieve desired
147 success is a thorough, systematic and clinically linked biological analysis of patient samples but
148 disease rarity and tissue processing challenges such as logistics and infrastructure have contributed to
149 a lack of relevant samples for clinical care and research. There is a need for a Europe-wide framework
150 to be implemented for the adequate and minimal sampling, processing, storage and analysis of patient
151 samples. Two international panels of scientists, clinicians and patient and parent advocates have
152 formed the Fight Osteosarcoma Through European Research (FOSTER) consortium and the Euro
153 Ewing Consortium (EEC). The consortia shared their expertise and institutional practices to formulate
154 new guidelines. We report new reference standards for adequate and minimally required sampling
155 (time points, diagnostic samples, liquid biopsy tubes), handling and biobanking to enable advanced
156 biological studies in bone sarcoma. We describe standards for analysis and annotation to drive
157 collaboration and data harmonisation with practical, legal and ethical considerations. This position
158 paper provides comprehensive guidelines that should become the new standards of care that will
159 accelerate scientific progress, promote collaboration and improve outcomes.

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168 INTRODUCTION

169 Osteosarcoma and Ewing sarcoma are malignant bone tumours affecting ~1,800 people annually in
170 Europe ¹. Despite continuous efforts and the investigation and intensification of treatment modalities,
171 the prognosis for patients is poor when compared to other cancers ^{2,3}. Repeated attempts by large
172 international cooperative groups to improve outcomes through randomised clinical trials have not led
173 to survival improvement in osteosarcoma ⁴⁻¹¹ and brought only modest benefits in Ewing sarcoma ¹²⁻
174 ¹⁸. A lack of available high-quality biological samples for omics (e.g. genome-wide profiling)
175 assessments has meant that we still have poor understanding of the molecular basis of observed
176 heterogeneous clinical phenotypes and mechanisms of chemoresistance and metastasis. Acquisition of
177 snap frozen and fresh tissue is recommended in international clinical guidelines ¹⁹⁻²², but is frequently
178 not achieved and the absence of standardised procedures for sampling has hampered compliance.

179
180 Two international panels of scientists, clinicians and patient and parent advocates formed the Fight
181 Osteosarcoma Through European Research (FOSTER) consortium (www.fosterconsortium.org) and
182 the Euro Ewing Consortium (EEC) ([https://www.ucl.ac.uk/cancer/research/centres-and-](https://www.ucl.ac.uk/cancer/research/centres-and-networks/euro-ewing-consortium/euro-ewing-consortium)
183 [networks/euro-ewing-consortium/euro-ewing-consortium](https://www.ucl.ac.uk/cancer/research/centres-and-networks/euro-ewing-consortium/euro-ewing-consortium)) to promote European collaboration and to
184 accelerate clinical and scientific progress. The consortia have already delivered benefits by bringing
185 together multiple – previously disparate – national clinical trial groups and scientists to develop and
186 deliver collaborative trial protocols ^{14,17,23,24}, share samples ^{25,26} and expertise ²⁷ to perform
187 collaborative research. A major goal of both consortia is the refinement and intensification of
188 translational research. Systematic acquisition of high-quality biological samples from children and
189 adults across multiple sites with associated clinical metadata should enable the identification and
190 characterisation of disease subgroups and tumour and germline genetic, biological, immunological
191 and cellular environmental factors that can be used for the stratification of disease subgroup-specific
192 therapies.

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194 This position paper complements international clinical guidelines and provides comprehensive
195 procedures for the adequate minimal sampling, handling and storage of bone sarcoma samples that

196 should be adopted across European centres. Although this statement has been drafted by the
197 osteosarcoma and Ewing sarcoma communities, the principles discussed apply equally to other bone
198 sarcoma histotypes and perhaps other cancers where a lack of samples hinders translational research
199 and clinical progress.

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201 **UNMET CHALLENGES IN OSTEOSARCOMA AND EWING SARCOMA BIOLOGY**

202 Key features of osteosarcoma biology include *in utero* loss-of-imprinting at chr.14q32^{28,29}, postnatal
203 *TP53* loss-of-function³⁰ (or possibly mutant gain-of-function³¹) and complex genome rearrangements
204 via chromoplexy and chromothripsis (^{32,33} and Valle-Inclan JE, Noon SD, Trevers K, Elrick H, Tanguy M,
205 Butters T, et al: Mechanisms underpinning osteosarcoma genome complexity and evolution.
206 bioRxiv:2023.12.29.573403, 2023). Specific molecular alterations in some cases include *MYC*
207 amplification³⁴, *RBI* deletion and mutation and a ‘*BRCAness*’ phenotype³⁵. Ewing sarcoma cells are
208 characterised by gain-of-function gene rearrangements between *FET* (*FUS*, *EWSR1*, *TAF15*) RNA
209 binding proteins and *ETS* (*FLI1*, *ERG*, *FEV*) transcription factors, most commonly *EWSR1::FLI1*³⁶.
210 The *FET::ETS* fusions encode oncogenic chimeric transcription factors with neomorphic features that
211 reprogramme the transcriptome³⁷, binding to GGAA microsatellites that become neoenhancers^{38,39},
212 which leads to ectopic gene expression and tumour development. Additional *STAG2* and *TP53*
213 cooperative mutations are associated with poorer survival⁴⁰⁻⁴³.

214

215 Although the key driver mutations and recurrent alterations present in a subset of cases have been
216 identified in both tumours, fragmented data from multiple small series and a lack of sufficient and
217 appropriate solid and liquid tissue biopsies have hindered the development of molecular
218 classifications and risk stratifications. Current and recent European clinical trials in Ewing sarcoma
219 (ISRCTN92192408, ISRCTN36453794, NCT00987636) have collected prospective liquid biopsies
220 and accessed clinical diagnostic tissue samples to validate previously reported prognostic biomarkers,
221 but none include specific molecular analysis of pre- and post-treatment tumour samples and clinical
222 trials are not representative of all patient groups. For osteosarcoma, there have been no large

223 prospective clinical trials since the closure of the EURAMOS-1 study and clinical trial samples do not
224 inform individual patient treatment decisions. A culture of more universal prospective tissue
225 collection is needed.

226

227 **‘REPRESENTATIVENESS’ OF CURRENT RESEARCH MODELS**

228 Preclinical models are a central component of translational research. Model systems such as patient-
229 derived cell lines, ex vivo engineered models ⁴⁴⁻⁴⁶ and spheroids / tumoroids ^{47,48}, in addition to in vivo
230 rodent (e.g. mice, rats), non-rodent (e.g. canine, zebrafish, xenopus) ⁴⁹ and chicken chorioallantoic
231 membrane ^{50,51} models, allow researchers to mimic bone sarcoma including its genetics and molecular
232 biology, local microenvironment, systemic dissemination and drug response. Most bone sarcoma
233 deaths occur because of the emergence of drug-resistant lung, bone and/or bone marrow metastases.
234 Orthotopic and patient-derived xenograft ⁵² and engineered mouse models ⁵³ recapitulating
235 disseminated disease are essential. Sampling paired treatment-naive and relapsed material is critical
236 for the development of relevant models to avoid unfavourable scenarios where preclinical drug
237 efficacy data generated using less relevant models appear promising ⁵⁴⁻⁵⁶ but the subsequent clinical
238 trials show no patient benefit ^{23,57,58}.

239

240 Historical cell lines, recent patient-derived cells and orthotopic xenograft mouse models have been
241 developed for osteosarcoma ^{52,53,59-63} and Ewing sarcoma ^{53,56,64,65}, but they typically over-represent the
242 higher-risk end of the disease spectrum. The Innovative Therapies for Children with Cancer (ITCC)
243 consortium has generated patient-derived xenografts for *in vivo* compound testing from children with
244 relapsed disease and includes some bone sarcoma models ⁶⁶⁻⁶⁸, but more representative and accessible
245 patient-derived cell lines, xenograft and genetically engineered autograft models that allow
246 simultaneous examination of the tumour, immune, extracellular and structural microenvironment are
247 needed ⁶⁹⁻⁷².

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249 **ACCESS TO NOVEL THERAPIES**

250 There is limited commercial incentive for the development of novel therapies for bone sarcoma. The
251 European Medicines Agency (EMA) implemented the European Union (EU) Paediatric Regulation in
252 2006, requiring the investigation of new therapies in children before marketing authorisation was
253 granted. A waiver system in the initial legislation was modified in 2015 ⁷³, strengthening the legal
254 requirement to investigate all therapies with a relevant mechanism of action for childhood cancer.
255 United States (US) Congress approval of the Research to Accelerate Cures and Equity for Children
256 Act ('RACE Act') enacted in 2020 gave the Food and Drug Administration (FDA) powers to mandate
257 paediatric clinical trials for new oncology drugs with a molecular target relevant to childhood cancers.
258 There is considerable alignment between the EMA and FDA and this concerted regulatory approach
259 has and will lead to greater opportunities for access to novel targeted therapies in children.

260

261 Although peaking in incidence in the 2nd and 3rd decades and occurring in older adults as well as
262 children, osteosarcoma and Ewing sarcoma are frequently considered 'paediatric' cancers and are
263 represented in early phase paediatric drug trials. The regulatory coordination between the EMA and
264 FDA presents an opportunity to utilise the inclusion of patients with osteosarcoma and Ewing
265 sarcoma in early phase trials to study drug response and to develop predictive biomarkers. However,
266 the number of patients with bone sarcomas recruited to each early phase trial is typically small ^{27,74},
267 sampling is not standardised, correlative biomarker studies are typically published long after
268 conclusion of the trial, if at all, and together these factors have led to an extreme paucity of high-
269 quality predictive biomarker evidence relevant to bone sarcoma. IGF1R inhibitors in Ewing sarcoma
270 are an example of a failed opportunity to identify why only some patients responded to treatment.
271 Across multiple early phase trials, multiple agents and over 400 patients, IGFR1 inhibitors resulted in
272 response rates of 5-15%, including some sustained responses ⁷⁵⁻⁸⁰, but no predictive biomarkers were
273 identified. As a result, no patient enrichment was possible in the Children's Oncology Group
274 AEWS1221 study comparing standard interval compressed VDC/IE with or without ganitumab. There
275 was no significant difference in survival between the arms.

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277 **THE IMPORTANCE OF OPTIMISING SAMPLE COLLECTION**

278 Several factors have converged to limit translational progress in osteosarcoma and Ewing sarcoma
279 including recurrent molecular alterations not being validated, a consensus on molecular classification
280 being made, a burgeoning of preclinical models but with an over-emphasis on high-risk disease, a
281 relative paucity of models for some disease settings, limited access to samples, almost non-existent
282 validated information about predictive biomarkers of response to cytotoxic chemotherapy and
283 molecularly directed treatment plus poor recruitment to early phase trials. In particular, while the key
284 molecular drivers of osteosarcoma and Ewing sarcoma are relatively well understood, there is a
285 fundamental lack of understanding of how genetic and epigenetic modifiers and tumour-host
286 interactions affect disease progression and treatment response. This lack of understanding is largely
287 driven by the absence of comprehensive, serial, annotated tumour tissue, normal tissue stroma and
288 liquid biopsy. At the level of clinical trials and collaborative large-scale research, there is a need for
289 more, high-quality, tumour and normal tissue (solid and liquid) biopsies, ideally, serial biopsies to
290 facilitate research into the molecular drivers and inhibitors of treatment response. At the level of
291 individual patients, tissue acquisition needs to meet the needs of modern, multi-omic analysis to
292 monitor disease response and facilitate options for molecularly targeted, personalised medicine and
293 critically for osteosarcoma, to identify patients with underlying cancer predisposition syndromes.
294 Taking tyrosine kinase inhibitors (TKIs) as an example, several TKIs have shown promise as single
295 agents in osteosarcoma and Ewing sarcoma⁸¹⁻⁸⁶ but despite responses in up to 40% of patients, there
296 are as yet no validated predicted biomarkers and the TKI mechanism of action remains obscure.
297 Ongoing trials are evaluating combinations of TKIs with chemotherapy in front-line and relapse
298 settings (e.g. the INTER-EWING-1 and rEECur trials developed by the EEC and NCT05691478 in
299 the USA) and the FOSTER consortium was recently awarded ATTRACT funding to investigate the
300 TKI cabozantinib as 12-month maintenance therapy following first-line standard therapy in
301 osteosarcoma. All include sampling timepoints designed to investigate biomarkers predictive of TKI
302 response.

303

304 A decades-long limitation to resolving some of the challenges discussed above is that there are no
305 consistent or systematic Europe-wide practices for sample collection. Standard operating procedures

306 (SOPs) for biopsies and other sample types, storage and sharing are either absent or only developed at
307 local or national level. Exacerbating the problem is that there is little infrastructure and few dedicated
308 staff to obtain bone sarcoma biopsies for both clinical care and translational research, although recent
309 initiatives are working towards changing this landscape.

310
311 Across Europe, the stakeholders engaged in obtaining biopsy material have different practices. The
312 amount, quality and availability of viable tumour material is variable and frequently inadequate for
313 molecular analyses. Because of the lack of a framework for sampling, much tumour tissue research is
314 performed on postoperative, necrotic material obtained after induction therapy meaning there is
315 ‘tainted’ data and knowledge on tumorigenesis, clonal evolution, metastasis and experimental drug
316 response. There is evidence that the chromoplexy attribute of osteosarcoma results in dramatically
317 different genetic alterations in different regions of the same tumour ⁸⁷, making a strategic approach to
318 tissue biopsy critical to understanding patient-specific tumour biology and target actionability (Table
319 1).

320
321 We present consensus guidelines on the appropriate type and timing of tissue and liquid samples to
322 facilitate research for future patients and to inform the treatment and future surveillance of current
323 patients. Where this dedicated approach has taken place in other cancers, for example melanoma, the
324 10-year survival rate has improved from ~10% ⁸⁸ to ~56% ⁸⁹ because high-quality samples are made
325 available for routine testing of the *BRAF* gene, which dictates first-line immunotherapy decision.
326 Cytotoxic chemotherapy is now disregarded as first-line therapy in melanoma.

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328 **GUIDELINES AND RECOMMENDATIONS**

329 **Introduction**

330 Cooperative effort from all involved disciplines is required. Routinely obtained written informed
331 consent, collection and storage of patient material for advanced biological studies is recommended in
332 international clinical guidelines ^{19,20,22} but non-compliance exists because of a lack of standard
333 procedures for biological sampling. Our position, complementing the clinical guidelines, is that all

334 patients with bone sarcoma should have snap frozen and fresh tissue samples (in addition to the
335 conventional diagnostic samples) taken at diagnosis, surgery and relapse regardless of their inclusion
336 in research initiatives or clinical trials.

337

338 Biopsies should be performed at specialist bone sarcoma units^{90,91}. Within research groups, clear
339 definitions of the sample types and relevant SOPs should be used. Solutions for ethical, legal and
340 practical issues should be widely shared. To maximise the advantages of sample collection, to obtain a
341 comprehensive biological understanding of bone sarcoma and host-related factors, different sample
342 types at sequential stages of the clinical pathway should be collected (Figure 1; Table 2). To enhance
343 fundamental understanding of bone sarcoma clonal evolution and chemoresistance, tumour tissue
344 collection at relapse and autopsy (e.g. PEACE study, NCT03004755) is essential. Metastases often
345 comprise different genetics to the original primary tumour so sampling metastatic lesions is
346 recommended to ensure that the maximal amount of biological information is collected.

347

348 **Diagnostic biopsy**

349 Treatment-naive core or open biopsies should be obtained from suspected bone sarcoma cases at
350 sarcoma specialist centres with the infrastructure to take, process and store (or send to a centralised
351 national centre) snap frozen and fresh tissue in addition to the biological material placed in formalin.
352 Fine needle aspiration is not adequate. Biopsies and their position should be determined at a
353 multidisciplinary team meeting with discussion on what the suspected lesion is expected to be, which
354 tumour zones the biopsies should be taken from and by which approach to avoid unnecessary
355 contamination. The procedure should be performed by a musculoskeletal or interventional radiologist
356 experienced in the diagnosis of bone tumours or by a specialist surgeon and reported in line with the
357 International Collaboration on Cancer Reporting²⁰ (<https://www.iccr-cancer.org/>). The biopsy tract
358 should be considered contaminated and resected en bloc during local therapy or be included in the
359 radiotherapy field to minimise the risk of local recurrence^{19,21,92-94}. The surgeon who will perform the
360 tumour resection should be involved in defining the optimal approach for the biopsy. The biopsy tract
361 is preferably marked and described according to compartmental anatomy^{92,95,96}. In many cases,

362 image-guided percutaneous biopsy using 8-, 11- or 14-gauge needles represents a well-established
363 alternative to open biopsy in terms of safety and diagnostic results ^{21,92-94}. Advantages and
364 (contra)indications have been described for both procedures ^{92,97-100}.

365
366 Sampling focused purely on histological diagnosis, usually from decalcified FFPE tissue, does not
367 consider the developing prognostic technologies that require snap frozen and/or fresh tissue that are
368 becoming standards of care, for example, the macrophage expression phenotype in osteosarcoma ¹⁰¹.
369 The equivalent of three 11- or 14-gauge needle biopsy samples have typically been sufficient to
370 provide diagnostic yield ¹⁰² when paired with conventional histology. Our position is that where a core
371 biopsy is performed, five samples should be collected where possible, of which at least one must be
372 snap frozen (Table 2). The 4th and 5th sample should be designated for research but can be used for
373 diagnostic purposes where a diagnosis could not be made using the FFPE samples. In many cases, the
374 4th and 5th sample will also be snap frozen and stored but depending on active research studies, one or
375 both could be formalin-fixed for use in spatial transcriptomics or used fresh for the isolation of live
376 tumour cells for cell line generation, organoid development and/or engraftment into
377 immunocompromised animals (Table 2). For open biopsy, a minimum of 1 cm³ of tissue cut into
378 multiple 0.2 cm³ sections is recommended. Where there are detectable oligometastases at
379 presentation, consideration should be given to obtaining metastatic tissue at the time of the biopsy.
380 For reference, recent Children's Oncology Group guidance advocates up to 20 core biopsies for bone
381 sarcomas with a soft tissue component (or up to 7 core biopsies where there is no soft tissue) plus up
382 to 3 cores of underlying osteoid ¹⁰³.

383

384 **Primary tumour resection and metastasectomy**

385 There are three surgical specimens where resection serves as both performing standard of care and
386 obtaining research samples: (i) primary tumour, (ii) matched adjacent normal tissue and (iii)
387 metastatic lesions. Samples should be prioritised by the pathologist to collect, depending on the
388 availability of biobanking and specific research initiatives: (1) FFPE as the standard of care and

389 neoadjuvant chemotherapy assessment, (2) snap frozen and stored, (3) fresh and placed into an RNA-
390 preserving medium, (4) fresh and placed into a culture-compatible medium (Table 2).

391

392 **Relapsed disease**

393 Samples from relapsed disease are particularly valuable if they can be paired with tissue from the
394 primary diagnosis. As most bone sarcoma recurrences develop early and there is usually little doubt
395 about the diagnosis, pre-treatment biopsy material is scarcer than at initial diagnosis. Given the poor
396 outcomes of relapsed disease and the limited treatment options, consideration should be given to
397 obtaining snap frozen and fresh and/or fixed tumour tissue at recurrence. These samples should be
398 appropriately processed for omics assessment, other research or biobanking. Irrespective of whether
399 there are currently recruiting and/or routinely commissioned omics initiatives available at the time of
400 recurrence, relapsed tissue is highly valuable if stored for future assessment.

401

402 **Blood samples**

403 Blood samples should be obtained at (i) diagnosis, (ii) before and after surgery and (iii) at follow-up.
404 Blood can be used as a liquid biopsy for the identification of circulating tumour DNA (ctDNA) and
405 RNA (ctRNA), circulating cell-free DNA (cfDNA) and circulating tumour cells (CTCs). For specific
406 diagnostic, monitoring and biomarker studies, urine and other body tissues (e.g. tears, hair) may be
407 collected. Blood samples should be processed according to the relevant study, for example, CTC
408 studies to be collected in cell-free Streck, PAXgene or EDTA blood collection tubes (BCTs) and
409 processed immediately. Streck and PAXgene both have BCTs specifically designed for ctDNA and
410 ctRNA capture. EDTA tubes can be used for either analytes, proteins or live cells. There are pros and
411 cons to each BCT related to the need for immediate versus delayed processing, plasma volume yield
412 and transport and storage costs. There is no consensus between European centres on which, if any, is
413 best overall. We recommend that EDTA is used as a minimum for storage as these BCTs enable most
414 analyses. But other more specific BCTs can be used according to research studies taking place at the
415 time of collection. Blood samples may be key to detect micrometastases as well as allowing for the
416 analysis of metastatic tumour-derived DNA, RNA (including microRNA) or proteins in circulation.

417

418 **Technical considerations**

419 Technical aspects of collection and storage need to be considered to obtain minimum amounts of
420 high-quality samples (Table 3), which may require a fundamental change in clinical practice in
421 individual centres. Radiologists, surgeons and pathologists have critical roles in the collection of
422 adequate samples for histological and molecular diagnostics and for translational research. The
423 biopsy, operative and histology procedures need to allow sufficient time to be devoted to sample
424 collection and processing. These procedures should be appropriately funded. If diagnostic centres are
425 unable to adequately process and store relevant material, consideration should be given by national
426 bodies to restrict diagnostic biopsies to centres with adequate infrastructure or establish regulated
427 delivery channels to central repositories.

428

429 **Standard operating procedures (SOPs)**

430 SOPs for tissue processing should be implemented by designated staff other than the radiologist or
431 surgeon because the tissue needs to be processed at the same time as the procedure being performed,
432 which requires the full attention of the radiologist or surgeon. After collection, material allocated by
433 the pathologist for diagnostic procedures will be processed as standard. Samples to be frozen should
434 be transferred to sterile vials and immediately snap frozen and stored in -80 °C freezers or in liquid
435 nitrogen. Fresh samples for cell and organoid cultures or animal engraftment need to be placed under
436 sterile conditions into appropriate vials with a culture-compatible medium. The logistics and reagents
437 may require pre-planning with the research group for material transfer to the laboratory within 24 h.

438

439 **Infrastructure and personnel**

440 Sampling requires a team effort. Some centres will require changes to current care pathways, for
441 example, automatic reminders to collect samples and duplicating processes so the biobank sample
442 pathway is parallel with the pathology sample pathway. The radiologist's and surgeon's focus will be
443 on the clinical procedure so it is important to establish a tissue processing pipeline as an
444 interdisciplinary effort and adapt it to local conditions, which may include oncology, pathology,

445 biobanking and theatre staff. For SOPs to work, theatre staff must be well informed, prepared and
446 adequately resourced to undertake the extra work. All personnel involved should recognise that tissue
447 processing for research is pertinent to future patients being cured. Understanding the importance of
448 their new role in tissue sampling could increase personnel efficiency and reliability.

449

450 **PATIENT AND PUBLIC SUPPORT**

451 Patients and their families overwhelmingly support research sample donation surplus to diagnostic
452 requirement. FOSTER together with the Sarcoma Patient Advocacy Global Network (SPAGN) have
453 undertaken an international survey. The survey includes questions on diagnosis, treatment and
454 survivorship experiences, plus assessment of patient and family priorities for future research. Four
455 questions are specific to sample donation. As of 2 February 2024, there were 372 combined
456 osteosarcoma and Ewing sarcoma respondents (n=234 osteosarcoma, median age 16 y; n=138 Ewing
457 sarcoma, median age 14 y). Just over half of respondents with Ewing sarcoma (52.2%) and less than
458 half with osteosarcoma (46.6%) were asked to donate research samples (Table 4). Of those asked,
459 97% consented to donate (Table 4). For the half of respondents who were not asked, almost two-thirds
460 reported that they would like to have been asked (Table 4).

461

462 **ETHICAL, LEGAL, PRIVACY AND PRACTICAL CONSIDERATIONS**

463 Responding to patient-led direction involves important ethical, legal, privacy and practical
464 consideration (Table 5). Patients or their families must provide written informed consent for the
465 collection, storage and use of research samples. Lawful protocols should be in place to ensure that
466 patient confidentiality and personally identifiable data are protected. Consent and protocols need to
467 navigate the range of legal frameworks of different European nations. Age-appropriate information
468 sheets for patients and their guardians must explain the purpose of the planned tissue storage and/or
469 research, the recipients of the material (either now or in future) and the use of pseudonymised clinical
470 data prior to providing forms for informed consent. Pairing sample data with pseudonymised clinical
471 data including treatment and imaging findings and where the law allows, explicit linkage to regional

472 and national cancer registries, should be possible. Ethical approval from international, national or
473 local authorities to study samples previously collected should be obtained.

474

475 Advantages of centralised versus decentralised (virtual) tumour banking and procedures to check for
476 appropriate tissue representation for interpretable biological results should be considered. Whether
477 sample availability should be defined as a mandatory inclusion criterion for patients going into
478 clinical trials should be evaluated by regulatory bodies and ethics panels. For clinical trials,
479 responsibilities of trial coordinators and local centres should be defined and adapted to applicable
480 laws and regulations. Adequate coverage of the local costs and shipment of samples by research
481 grants or national initiatives can help to facilitate the compliance of local institutions, particularly
482 where there are financial challenges faced by sample collection units. However, in some cases,
483 financial constraints will prevent the collection of samples for unspecified research. Reusable tumour
484 box devices can facilitate the shipment of frozen and unfrozen material. Practical aspects of exchange
485 (including transborder) and the use of material should be defined by material transfer agreements
486 (MTAs) between research centres.

487

488 **BIOBANKING**

489 Biological material can be stored centrally by an academic tissue bank with software systems
490 allowing for maximal up-to-date information about the stored materials. The materials can also be
491 stored in local tumour banking facilities and later shipped in batches, as required, for use in further
492 analyses. Both centralised and decentralised material storage allow for their use in big data analyses
493 with bioinformatics support. Regardless of storage location (e.g. accredited laboratories with alarm
494 monitoring versus research lab freezers), proper evaluation by experienced bone sarcoma pathologists
495 should ensure appropriate tissue representation before being used in specific projects. Biological
496 material storage in aliquots allows for the tissue to be used for multiple research projects. Within
497 existing legal frameworks of some European countries it has been possible in some clinical units to
498 store fresh and snap frozen material from the biopsy before a diagnosis is obtained for a limited time
499 prior to explicit patient consent for biobanking¹⁰⁴. This practice requires the appropriate infrastructure

500 to be in place at the time of the biopsy and some bureaucracy to ensure adequate record keeping.
501 Across most European centres, it is the tissue bank where the samples were collected that owns the
502 biological material. Tissue banks are typically non-specific repositories for all patient materials
503 collected at a local institution or within a region, or sometimes can be a study-specific biobank.

504
505 Transparent criteria for the regulation of access to larger material series by researchers from local
506 contributing institutions could positively influence the cooperation of local centres. MTAs and SOPs
507 for material shipment and adequate cost coverage (e.g. research grants) could further facilitate
508 cooperative tumour banking. It is also important to establish procedures for the coupling of tumour
509 material data to patient data. Genomic, transcriptomic, methylomic and metabolomic data from
510 tumour biopsies plus data from experiments on patient-derived cell cultures and xenografts should
511 ideally be stored in an international bone sarcoma registry together with comprehensive anonymous
512 clinical, radiological and pathological data. It is worth investing in the collection of large amounts of
513 retrospective clinical data regarding baseline characteristics, treatment and survival from multiple
514 international groups and to correlate these data with the analysis of genomic and epigenomic data
515 from corresponding banked tumour samples. FOSTER, the EEC and clinical trial groups should
516 consider aspects of data collection and sample storage and discuss early in the planning phase of
517 collaborative projects so that specific national requirements and future projects linking datasets can be
518 implemented in a timely manner. Data sustainability beyond individual projects and connection of
519 data at overarching levels should be considered.

520

521 **CONCLUSION**

522 Tangible progress in bone sarcoma has been bottlenecked by insufficient biological assessment and
523 investigation, which in significant part has been caused by limitations in sample collection. Routine
524 collection of decalcified and formalin-fixed tissue for histological examination will not support
525 diagnostic and prognostic technologies that evolve from translational research, for example NGS, in
526 large part because fresh and snap frozen tissue is not routinely stored. The benefits of obtaining fresh
527 and snap frozen samples at biopsy exceed the risks of complications of taking more tissue. Changing

528 the process in which we collect biological samples and link patient data will lead to new molecular-
529 based standards of care as well as new targeted therapies with fewer side effects.

530

531 Metastatic sites are not routinely sampled. Liquid biopsies are not routine. Screening for germline
532 predisposition syndromes is not routine. The availability of properly sampled and stored biological
533 materials will confer multiple scientific and clinical advantages including allowing identification and
534 validation of new and reported prognostic factors and druggable targets. We need to ensure that
535 children, teenagers and young adults with bone sarcoma are not left behind while precision oncology
536 offers new treatment solutions for more common, typically older adult, cancers. Because paediatric
537 sarcomas are clinically and biologically highly distinct from adult cancers, precision medicine
538 approaches should be adapted to make the best use of samples that are as informative as possible.
539 Appropriate sample collection, storage and sharing can only be achieved successfully if all the
540 relevant steps are optimised at each local centre. Collection and storage procedures could be adapted
541 by local institutions to suit their individual structures, defined and assigned to dedicated individuals
542 who are specifically educated and trained. FOSTER, EEC and institutional researchers should actively
543 collaborate, share data, methods and samples and disseminate good practice. These approaches will
544 advance progress in bone sarcoma.

545

546 **AUTHORS' CONTRIBUTIONS**

547 This paper was mainly written by DG, RVE, ET, QCH, MGM and MN. Data from the sarcoma
548 experience survey were generated by PP. Authors are Young Investigators, Work Package and Nation
549 Leads who all contributed to the content of the paper from the perspective of the group that they
550 represent as well as commenting on general aspects of this Policy Review.

551

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563

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892 **FIGURE LEGEND**

893 **Figure 1: Overview of the sample types to be collected.** To maximise the advantages of sample

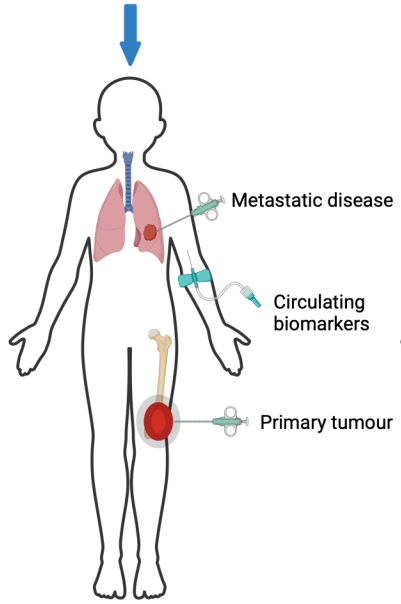
894 collection, in order to obtain a comprehensive biological understanding of bone sarcoma and host-

895 related factors, different sample types at sequential stages of the clinical pathway should be collected.

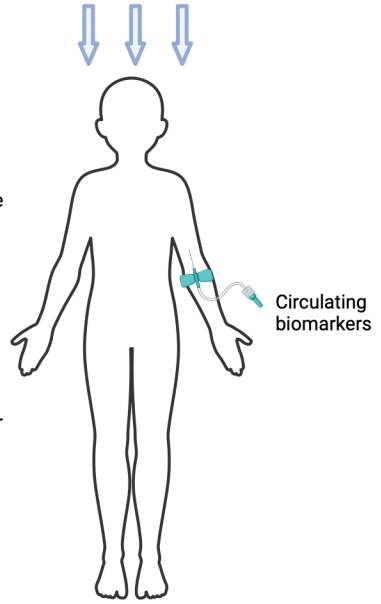
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Primary treatment

Standard of care sampling

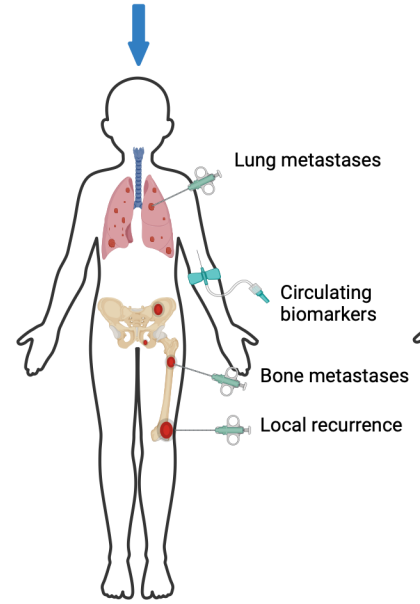


Serial sampling within a specific study



Recurrence

Standard of care sampling



Serial sampling within a specific study

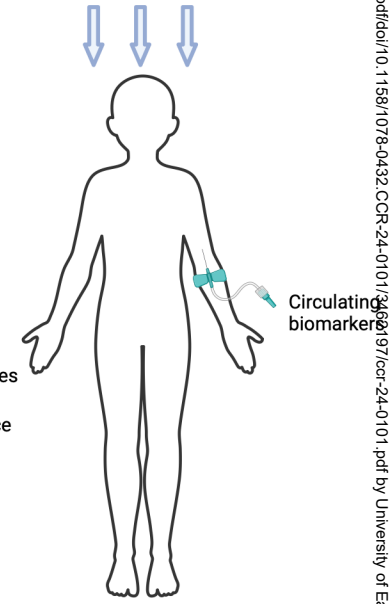


Table 1. Advantages of appropriately consented, collected and/or biobanked samples.

For individual patients

- Druggable target identification and screening for early phase trials
- Identification of germline predisposition syndromes
- Monitoring of minimal residual disease*
- Assignment to molecular strata*
- Therapeutic use for immuno-oncology approaches such as tumour vaccines*

For future research

- Identification and validation of molecular stratification
- Identification of mechanisms of pathogenesis, drivers of tumour growth and resistance mechanisms
- Analysis of biological drivers of relapse, particularly if paired diagnostic/relapse samples available
- Analysis of tumour microenvironment and immunological aspects
- Prognostic and predictive biomarker development and validation
- Validation of liquid biopsy methodologies and development of minimal residual disease biomarkers
- Identification and validation of SNVs associated with pharmacokinetic properties and treatment-induced early and late toxicities
- Establishment of representative preclinical models and patient-derived cell lines

For future research – particular benefits of prospective clinical trial samples

- Uniform sample processing, homogeneously treated patients
- Uniform clinical datasets within and between trials**
- Cross-validation of liquid biopsy, molecular classification, prognostic and predictive biomarkers between independent cohorts

* Assumes successful completion of ongoing research

** Aided by ongoing Pediatric Cancer Data Commons initiatives (<https://commons.cri.uchicago.edu/pcdc/>)

Table 2. Guidelines for sample collection ensuring diagnostic and translational research efficiency**Standard of care at: diagnosis, primary tumour resection, metastasectomy, recurrence**

	Processing	Purpose
Minimum essential 3-5 core biopsies using 8-, 11- or 14-gauge needles. OR larger cores divided into two or three pieces OR 1 cm ³ open biopsy cut into multiple 0.2 cm ³ pieces.	FFPE At least 1 core or tumour piece snap frozen in liquid N ₂ or immediately stored in -80 °C	Diagnostic Diagnostic & Research
Optimal 5-7 core biopsies using 8-, 11- or 14-gauge needles. OR larger cores divided into pieces OR 2 cm ³ (or 2 x 1 cm ³) open biopsy cut into multiple 0.2 cm ³ pieces plus normal tissue comparator.	Material to be snap frozen or fresh material used in ongoing research projects to develop PDXs, tumour organoids, primary cultures, etc.	Research
Optimal whole blood* in EDTA or other normal tissue for germline sequencing**	PBMCs, plasma, serum	Research

Samples for specific research studies and/or biobanking***

Live cells in a culture-compatible medium/organ transplant preservation solution	Tumour cells	Research
whole blood* in EDTA or PAXgene tubes	Circulating tumour cells	Research
whole blood* in EDTA or cell-free Streck tubes	Circulating tumour DNA, plasma, serum, PBMCs	Research
1-5 mL other biofluids	Saliva, urine	Research

Samples at death/autopsy

Oligometastases samples	Snap frozen in liquid N ₂ or immediately stored in -80 °C	Research
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*=procedures and volumes for children and adults in accordance with the WHO guidelines on drawing blood: best practices in phlebotomy. **=germline sequencing is not currently international standard of care but many European countries have ongoing standard of care NGS studies that include germline

sequencing. ***=blood samples may be taken serially during and after treatment where specific research projects are available

Table 3. SOPs to be considered in local institutions. Staff from all involved disciplines (e.g. interventional radiologists, surgeons, operating room staff, pathologists, paediatric and medical oncologists and research nurses) should be aware of the importance of the availability of adequate biological samples and define the practical steps of collection, storage and shipment of samples according to local structures.

- Obtain information and written informed consent from patients or their legal guardians.
 - Determining the amount and types of tissue, blood and other material to be collected.
 - Orthopaedic surgical considerations (frozen section, infiltration zone, margin material); freezing and fixation of maximal amounts of material.
 - Orthopaedic and pathological diagnosis and reference assessments.
 - Sending MRI data via digital route or anonymised and coded external drive.
 - Providing adequate short-term storage of tumour tissue and other samples.
 - Transferring materials to long-term storage or shipping samples according to SOPs.
 - Ensuring trial-specific requirements are met (e.g. tumour sections not sent to pathology for analysis but straight from the operating theatre to the research lab).
 - Supplying material for cell culture in specific sterile cell culture medium.
 - Filing documentation of collected materials per study in institution-specific lists or databases.
 - Confirming received materials at research institute.
 - Establishing procedure for prioritisation of pathology in case of sparse material.
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Table 4. The Patient and Parent Advocacy Group and Sarcoma Patient Advocacy Global Network international survey on sarcoma experiences. The survey has so far included 598 respondents with bone sarcoma experience, of which there are 234 with osteosarcoma and 138 with Ewing sarcoma. Questions were asked on diagnosis, treatment and survivorship as well as priorities for future research. Four questions were specific to research sampling.

Question	Osteosarcoma			Ewing sarcoma		
	Yes (%)	No (%)	Other (%)	Yes (%)	No (%)	Other (%)
Were you or your child/family asked to donate tissue samples for research?	109 (46.6%)	84 (35.9%)	41 (17.5%)	72 (52.2%)	39 (28.3%)	27 (19.6%)
If you or your child/family member were not asked to donate tissue, would you have liked to be asked?	82 (65.6%)	3 (2.4%)	40 (32%)	40 (60.6%)	2 (3%)	24 (36.4%)
If asked, did you or your child/family member agree to donate tissue?	106 (97.2%)	0 (0%)	3 (2.8%)	68 (94.4%)	0 (0%)	4 (5.6%)
If you or your child/family member consented, was tissue successfully collected/donated?	63 (59.4%)	0 (0%)	43 (40.6%)	42 (61.8%)	1 (1.5%)	25 (36.8%)

Table 5. Ethical, legal, privacy and practical aspects of sample storage, sharing and shipment between research groups.

- Age-appropriate information sheets must explain the purpose of the planned research, the recipients of the material and the use of anonymised or pseudonymised clinical data.
 - Coupling of tumour material data to patient data, including treatment and imaging findings.
 - Ethical approval and permissions from international, national or local authorities.
 - A monitoring system for available samples and for associated informed consents per local hospital.
 - Ownership issues relating to biological tissue and clinical data, which might be different between countries, should be considered.
 - Advantages of centralised versus decentralised (virtual) tumour banking and procedures to check for appropriate tissue representation for interpretable biological results should be considered.
 - Adequate coverage of the local costs and shipment of samples by research grants can facilitate the compliance of local institutions.
 - Integrated, reusable tumour box devices can facilitate the shipment of frozen and unfrozen materials.
 - Practical aspects of exchange and use of biological samples should be defined by MTAs between institutions.
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