

Thanks for a nice meeting in Malmö, it was very great to see everyone again. I hope to see you all in Stockholm next time!

This is a summary of the projects we discussed in Malmö.

### **Ongoing projects:**

**Epithelioid sarcoma, *Principal investigator F Mertens*:** Cases were reviewed at the Lund and Gothenburg meetings. I have received patient information from Trond/Tor but not from others. I'm attaching a new copy of the patient information scheme with this email, please fill out the data you have available and submit to me. For those who can submit tissue for genetic analysis, sections or DNA should be sent to Fredrik Mertens in Lund (details will follow in the end of this email). I know the Norwegian centers are currently processing a general ethical application. While no specific deadline has been set, the study will close before next summer, otherwise this study will never progress. Additional cases can be included as long as they have typical morphology and loss of INI1. If you have problems regarding ethical permission, funding or something else please let us know. We have received some funding for this project.

To do:

- i) send patient data to Felix Haglund ([felix.haglund@ki.se](mailto:felix.haglund@ki.se))
- ii) send available tissue/DNA to Fredrik Mertens for genomic array

**Low-grade fibromyxoid sarcoma / SEF, *Principal investigator Florian Puls*:** Cases were reviewed at the Gothenburg meeting. Florian will send out a review file for each center to fill out. Sections or DNA should be sent to Fredrik Mertens in Lund for analysis. Should we reserve time to review additional LGFS/SEF cases in Stockholm?

To do:

- i) send patient data to Florian Puls ([florian.puls@vgregion.se](mailto:florian.puls@vgregion.se))
- ii) send available tissue/DNA to Fredrik Mertens

### **New / potential projects:**

Before we decide upon which diagnosis/diagnoses to review in Stockholm, please let me know before the end of January how many cases you have of each diagnosis below. I will send a summary to everyone.

1. Intimal sarcomas (all primary cardiac sarcomas?)
2. Clear cell chondrosarcoma (Felix can finance sequencing)
3. Clear cell sarcomas
4. Osteblastoma / osteblastoma-like osteosarcoma

**MATERIAL FOR GENETIC ANALYSIS:** 10 micrometer sections with an area of 250 square mm. E.g. A tumor which measures 2,5 x 1,0 cm = 1 section; 1,3 x 1,0 = 2 sections). It is preferred that the first section from the block is not used to increase the quality of the material. Sections can be submitted in Eppendorf tubes (1 case per tube) and sent by regular mail. Alternatively, DNA can be prepared directly at the pathology department. It is not preferable to send whole tissue blocks.

Minutes made by Felix Haglund

Pehr Rissler  
Chairman