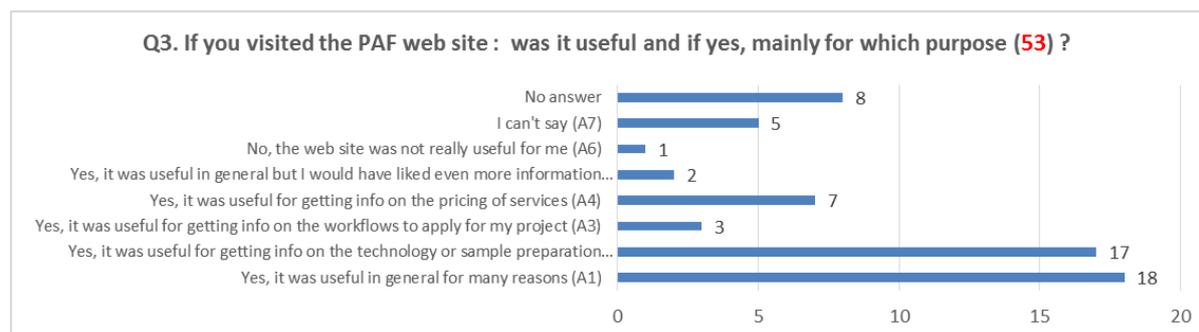
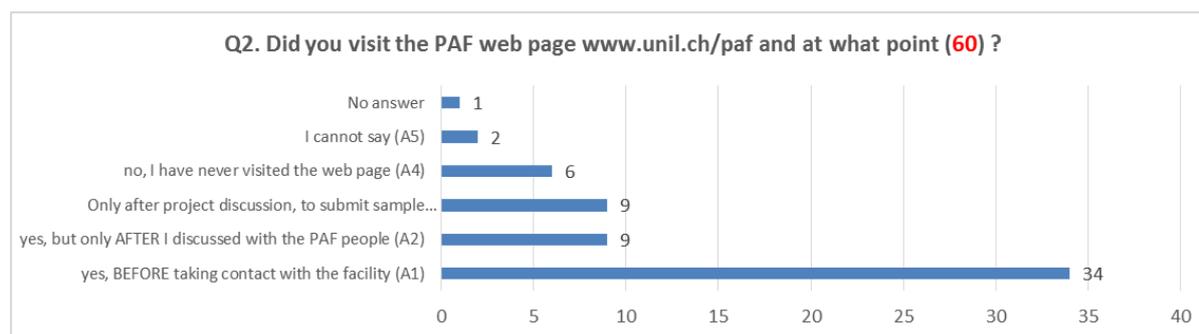
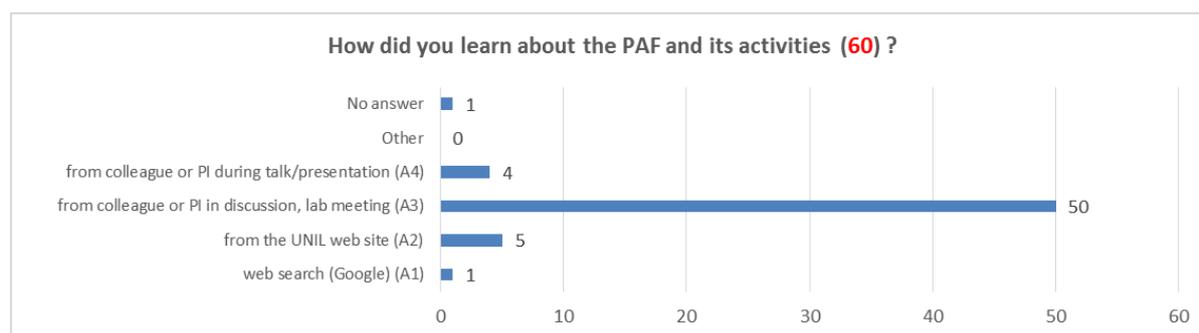


Results of the satisfaction survey 2015-2017

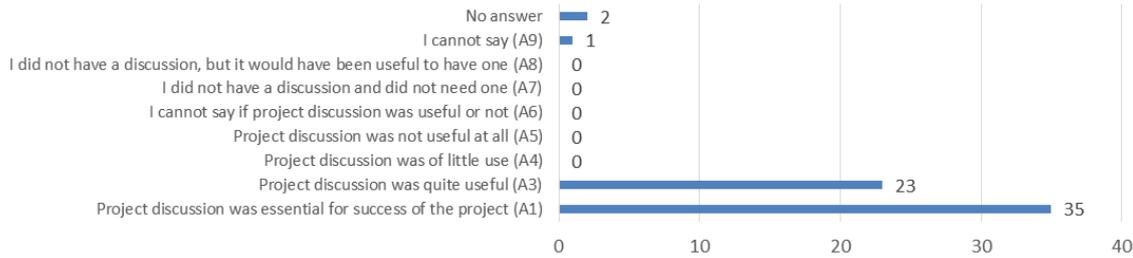
103 users of the platform between 2015 and June 2017 were contacted by e-mail; of these, **88** still had valid e-mail addresses (most of the 15 with invalid addresses had moved to a new position). Out of the 88 users that received the messages, **54 filled in the form completely (61.4%)**. The number of non-null answers is indicated in red in each graph.

Note : to ensure objectivity in handling , the survey was conducted and the results collected and analysed by a person external to our group.

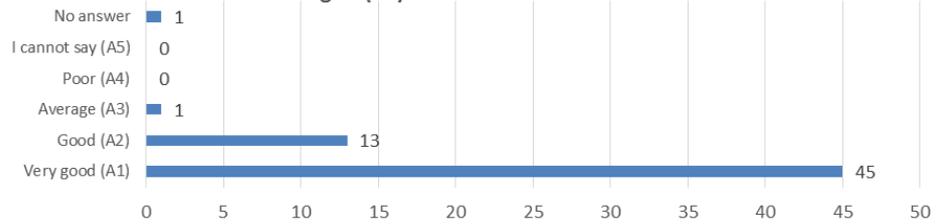
Note 2 : the voucher for free analyses was won by Laia Martinez Carreres, group Fajas.



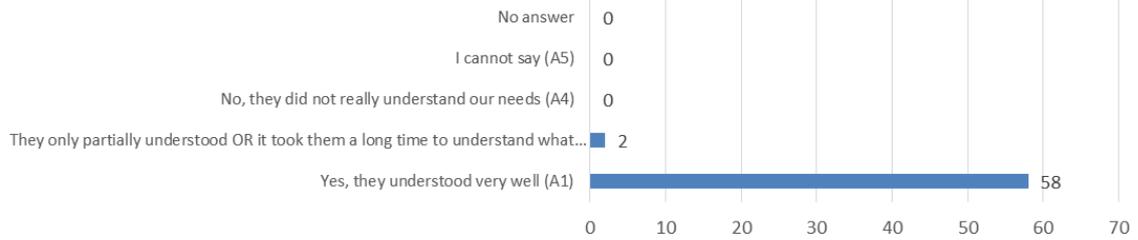
Q4. If you had an in-depth discussion of your project with PAF personnel, please rate this discussion (59).



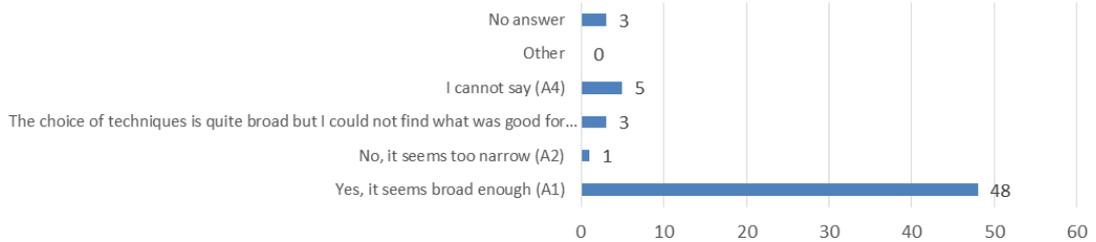
Q5. How do you rate the competence of the PAF personnel, concerning proteomics technologies (59) ?



Q6. Do you think that the PAF personnel was able to understand the specific needs of your project and suggest a suitable strategy (60) ?



Q7. Do you think that the palette of services offered by the PAF is broad enough (57) ?



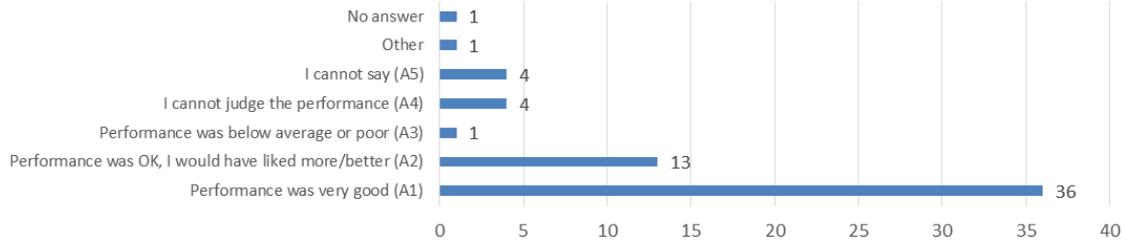
Q8. Targeting of services. Did you find the right service for you (59)?



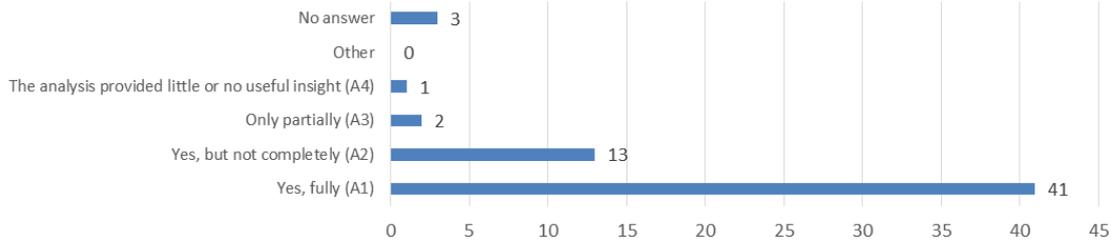
Q9. Quality of sample handling. Do you think that your samples were processed in an appropriate way as to get the maximum possible results (59) ?



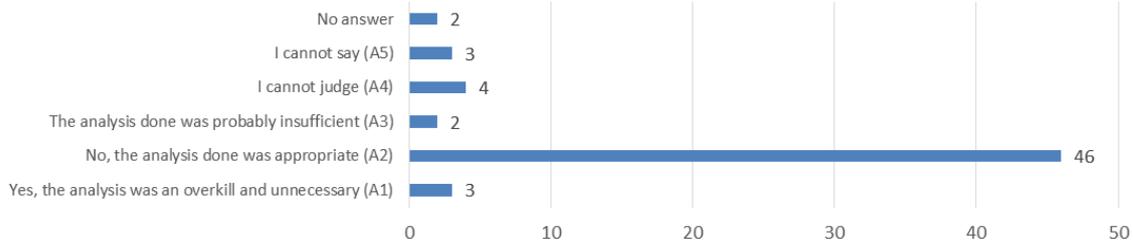
Q10. Analytical performance. If applicable how do you find the technical performance of the PAF MS analyses : were sensitivity and proteome coverage as you expected (59)



Q11. Analytical performance. Overall, did the PAF analysis answer the question(s) asked in the context of your project (57) ?



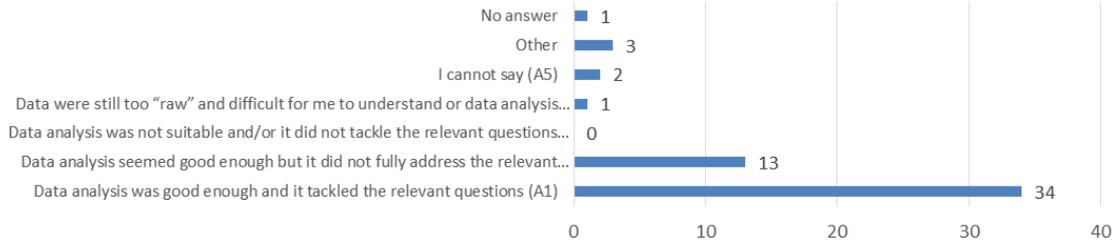
Q12. Was the analysis an overkill ? Do you have the feeling that the question relevant for your project could have been answered with a simpler, cheaper analysis (58) ?



Q13. Added value of analyses. Do you think that the proteomics data you obtained can be exploited/mined for other projects/questions or to generate (new) hypotheses (59) ?



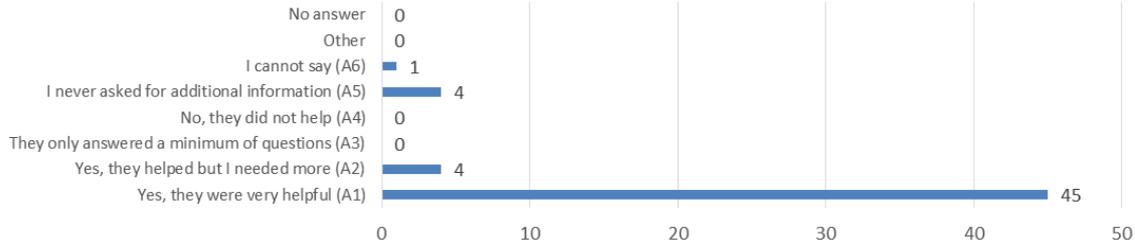
Q14. On the data analysis performed by the PAF. Was it sufficiently complete, and did it answer the questions relevant for the project (53).



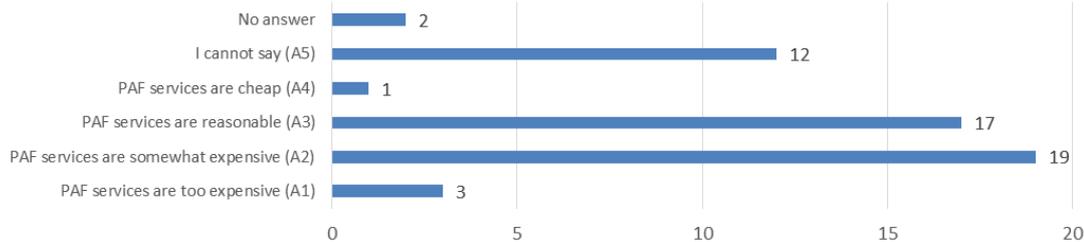
Q15. Results clarity For your specific project, were the results and the report received clear enough (54) ?



Q16. After the analysis, did you get sufficient support from the PAF personnel to discuss the results of the experiment, ask questions and solve data analysis issues (54) ?



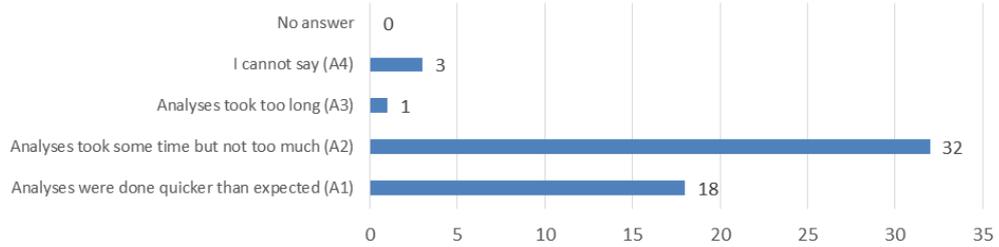
Q17. Please evaluate our service fees in absolute terms. (Please have a fair evaluation and if possible consider pricing of other tech platforms) (52)



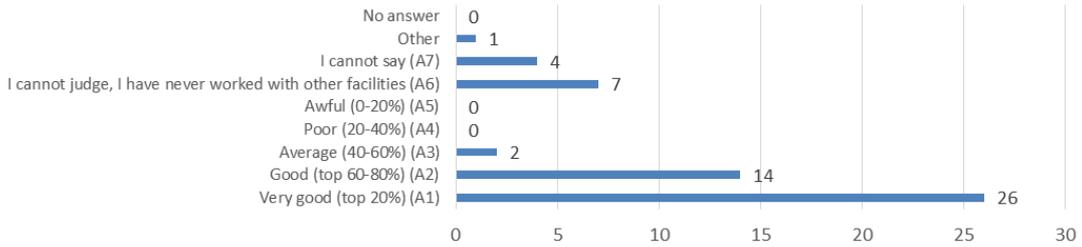
Q18. Price/performance ratio. Do you think that what you spent was too much or too little for the results obtained and the effort involved (50)?



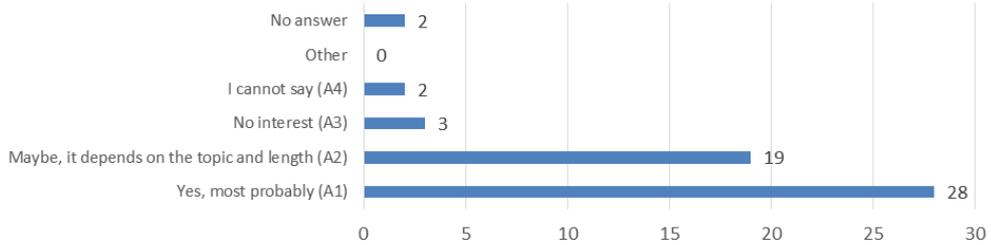
Q19. Processing times. Please evaluate the time from sample deposition to result generation. Please if possible try to consider the complexity of the experiments performed (54).



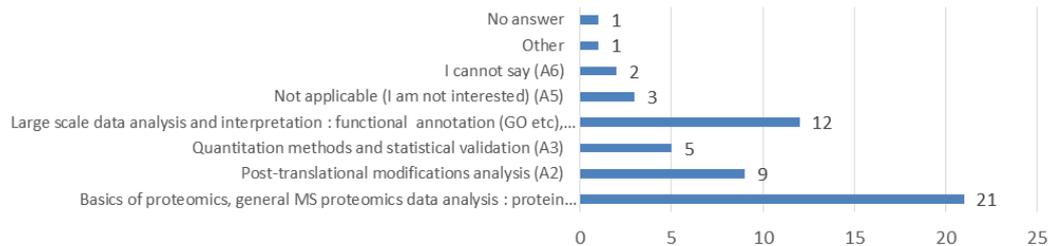
Q20. Global evaluation. Based on your experience, and comparing it with other proteomics facilities or research labs, how do you rate the PAF on a global scale of proteomics labs (54)



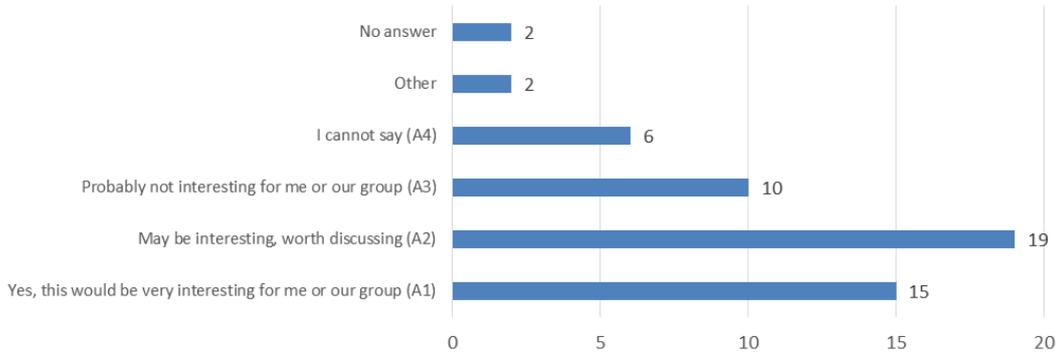
Q21. Offers of training. If the PAF would offer courses/workshops on proteomics technologies and data analysis, would you be interested to attend (52)?



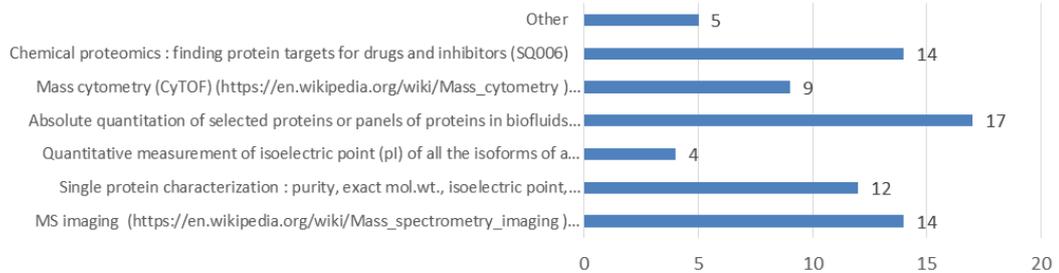
Q22. Training topics. Which topics of training courses would be more interesting for you (53) ?



Q23. "Advanced users" mode. We could consider to establish a mode of closer collaboration with strong users of proteomics technology. In this scenario, groups that have a major interest in proteomics experiments would send staff to the facility to be tr



Q24. Would you like to see other or new technologies implemented at the PAF as a service ? Please see some possibilities below (70)



From Q24, the suggestions for other techniques to implement were :

- i) I don't work with research anymore.
- ii) native MS
- iii) Protein design
- iv) Single cell western
- v) XL-MS

The last Question in the form (Q25) was a free feedback space (all responses given below)

Question 25

Any other comments/criticisms We appreciate any type of constructive input and/or criticism. Do not be afraid to write, this form is anonymous.

| | | |
|-----------|----|--------|
| Answer | 12 | 22.22% |
| No answer | 42 | 77.78% |

Excellent service. Very professional. Productive.

Following a seminar given by Prof. Matthias MANN from the Max-Planck Institute, it seems that the analysis we got from the PAF is not deep enough. It could be interested to get some fine tuned techniques his group developed.

I only want to say thanks for all your help.

I really appreciate the support of the PAF team from the beginning (writing) until the end (data analysis) of the project.

I wish you all the best in your research.

If you manage to shorten analysis time that would be really great

In general PAF does a good job.

My experiences with PAF have been joyful and very fruitful by large. Only seldom, at times, the time-gap between sample submission and result exceeds more than usual time. Other than that, I find the quality of service at PAF and the friendliness of staff very useful. My best regards.

No reason to complain.

Please note that I have requested the PAF assistance for a very unusual application. This did not require data analysis. However, the PAF has been instrumental in this project.

We had very successful collaboration with the PAF, one of the best proteomic facility I know

You guys are really excellent. I've never worked with a facility who were so competent, easy to work with and forward thinking as yours. I hope we can continue working together.