GIW2000 (11th Genome Informatics Workshop) は2000年12月18日と19日に、東京東京大学で開催され、約540名の参加がありました。今回から、そのポスターセッションとソフトウェアデモンストレーションが日本バイオインフォマティクス学会の年会として分離されています。今回のプログラム委員会は、米国ワシントン大学のKeith Dunker博士と北陸先端科学技術大学の理化学研究所の小長谷明彦博士が委員長を務められ、全部で20件の口頭発表が採択されました。これらの論文の要約はMEDLINEデータベースにも収められます。また、ポスターデモンストレーションが5件ありました。さらにセレラ・ジェノミクス社のGene W. Myers博士と東京大学の理化学研究所の横山茂之博士が最新の研究成果を招待講演で話してくださりました。なお今回、学会として、海外から参加して口頭発表を行い若手研究者の旅費の一部を援助することとし、2名の方が選ばれました。以下にその二人からのGIWレポートを掲載します。

GIW2001も同じ日程（12月18〜19日）、同じ場所で開催され、プログラム委員長には、Linsoon Wong博士と松田秀雄博士が就任される予定です。

Dr. Gene W. Myers

http://giw.ims.u-tokyo.ac.jp/giw2001/

Dr. Shigeyuki Yokoyama

■GIW Report from Christian Blaschke

Thanks to the generous travel award I received from the Japanese Society of Bioinformatics it was possible for me to attend to this very interesting workshop. My first impression was the good organization that was done at this event. The posters were presented all together in one part with enough pace to walk through, to get in contact with the authors and to discuss the work together.

Our own work is concentrated on text analysis applied to the results of expression experiments with DNA arrays. Therefore, it was very interesting to see that a considerable part of the posters was focused on the work with DNA arrays (around 20) and also some efforts in the field of text analysis could be seen.

The exhibition of the different hard and software companies was very informative. They demonstrated their new products and state of the art in professional bioinformatics. The only critique to this point may be that the exhibitors at an international conference should in general know English and be able to explain their products to a foreign person.

My experience with the oral presentations was also very positive. The microphones and the sound system and the lights were used in a very professional way. At the time of my own presentation I felt the help of the organizers with the microphones and the quality of the projector for the presentations was very high. The different chairmen did a very good job and most of the talks finished in their scheduled time.

The presentations came from a wide range of different fields. They included genome analysis, text analysis, DNA expression analysis, protein structure analysis and the simulation of genetic networks.

The program was very dense but there was enough time left to discuss the works in the breaks in a personal conversation with the authors. Concluding I have to say that this workshop was a very positive experience and it extended my knowledge in different fields of bioinformatics.

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