

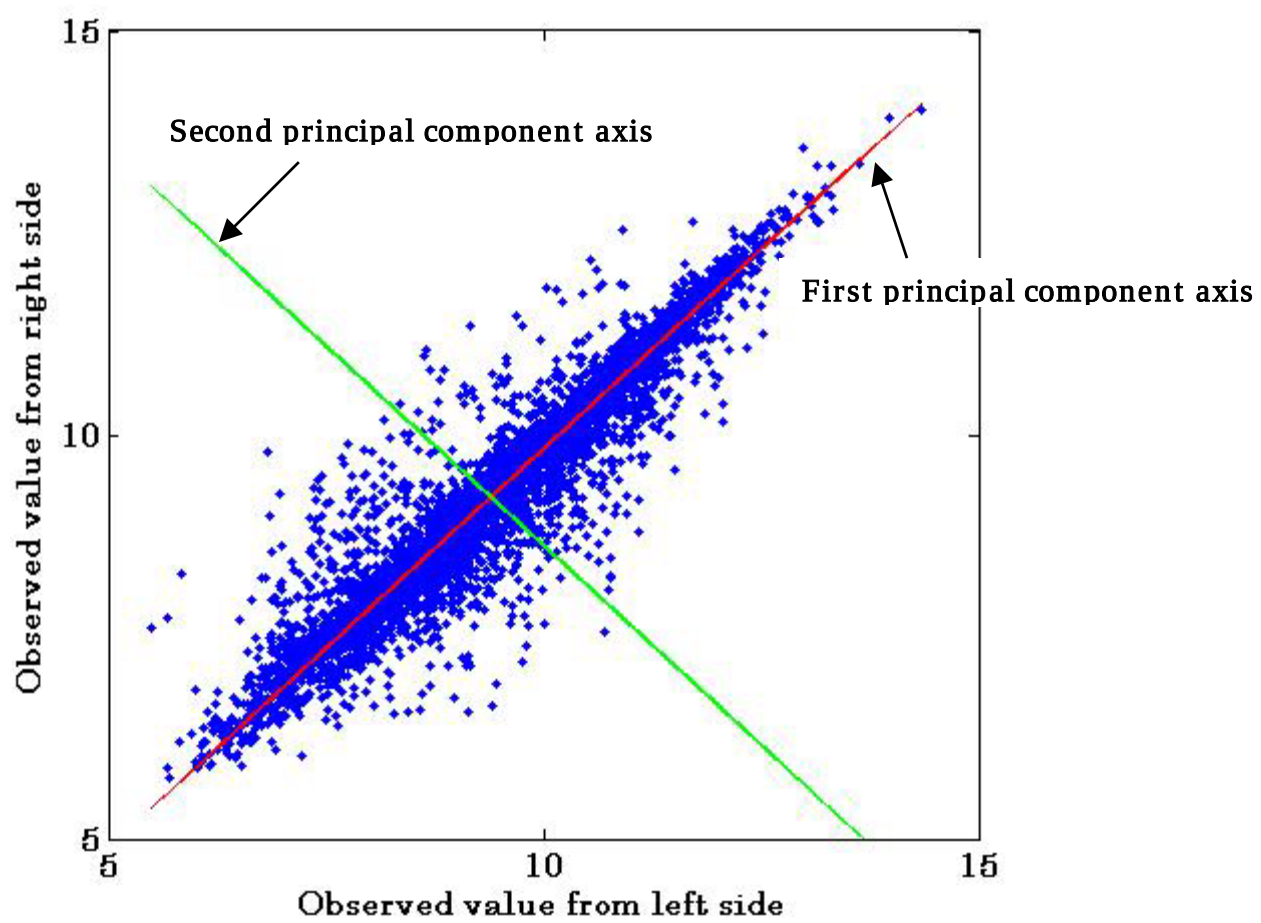


Fig.1.

M. Takeya et al. Breadth: 8.6 cm.

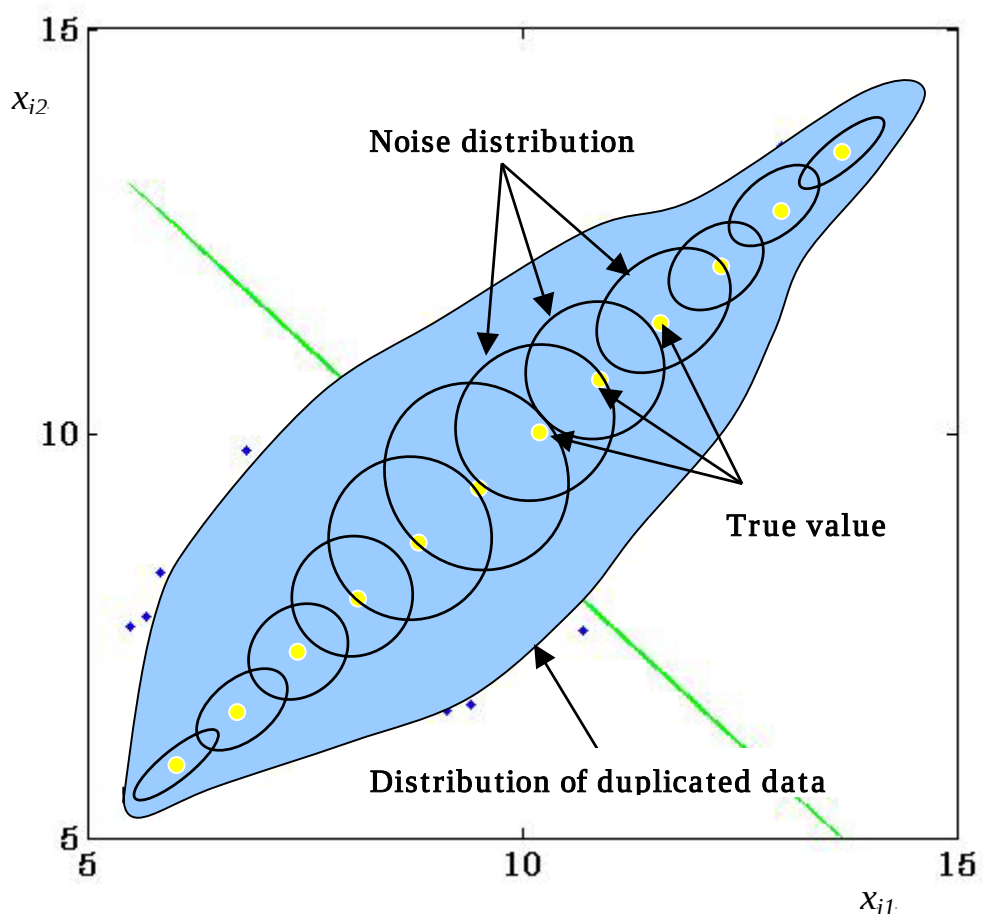


Fig. 2.

M. Takeya et al. Breadth: 8.6 cm.

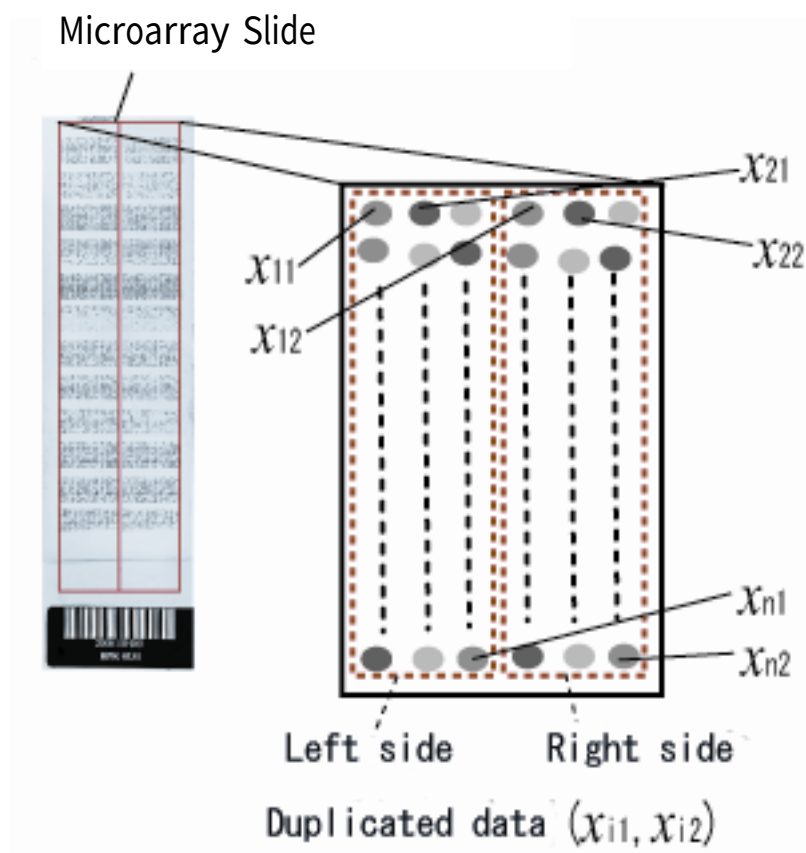


Fig. 3.

M. Takeya et al. Breadth: 8.6 cm.

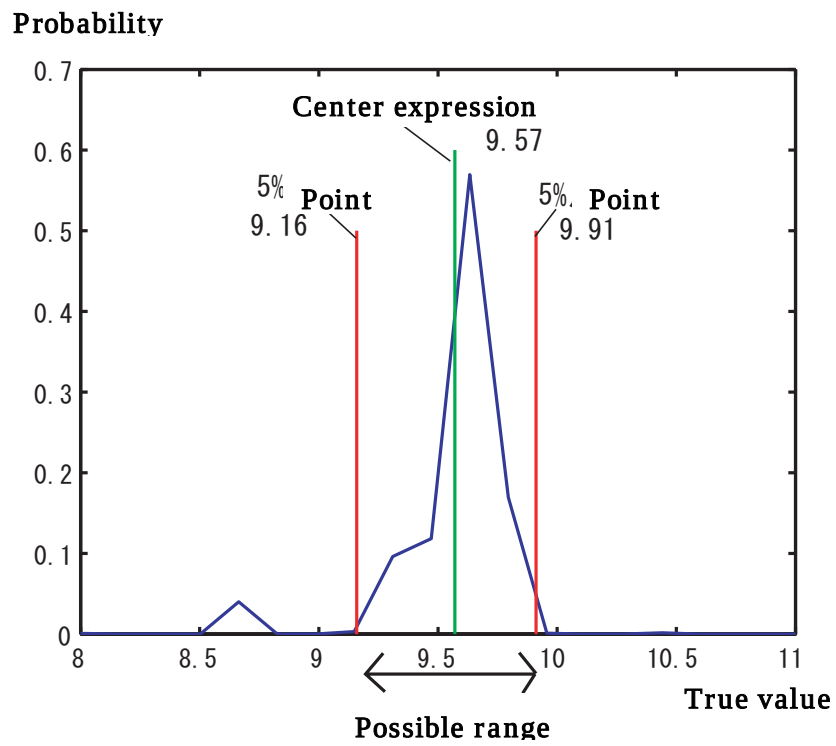


Fig. 4.

M. Takeya et al. Breadth: 8.6 cm.

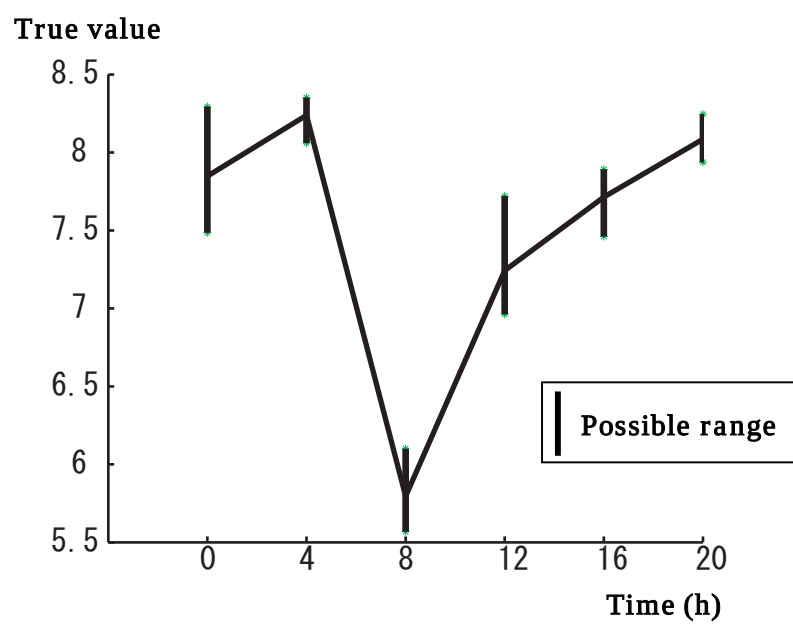


Fig. 5.

M. Takeya et al. Breadth: 8.6 cm.

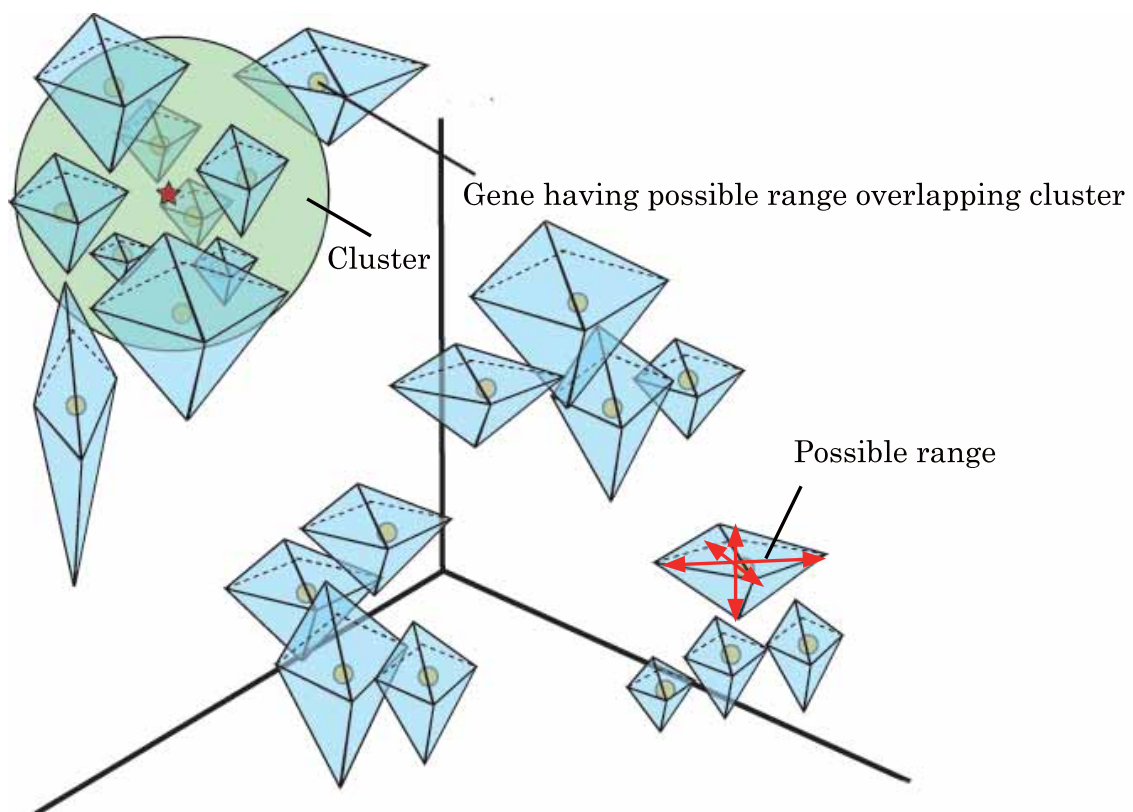


Fig. 6.

M. Takeya et al. Breadth: 8.6 cm.

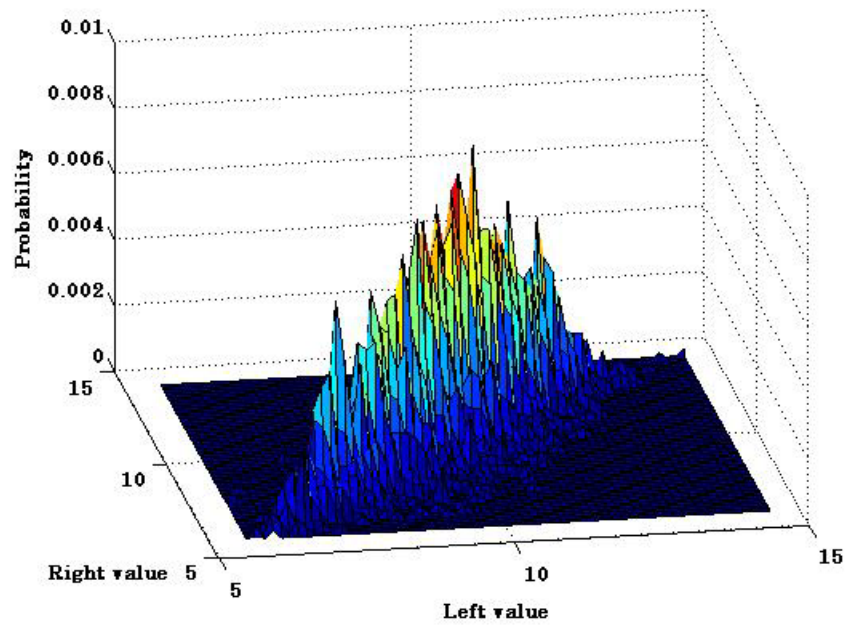


Fig. 7. (a)

M. Takeya et al. Breadth: 8.6 cm.

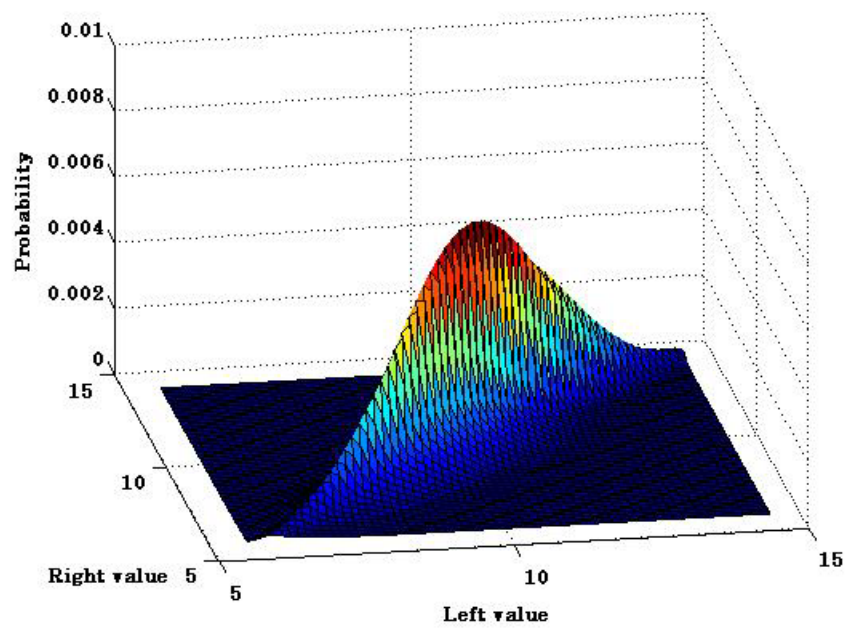
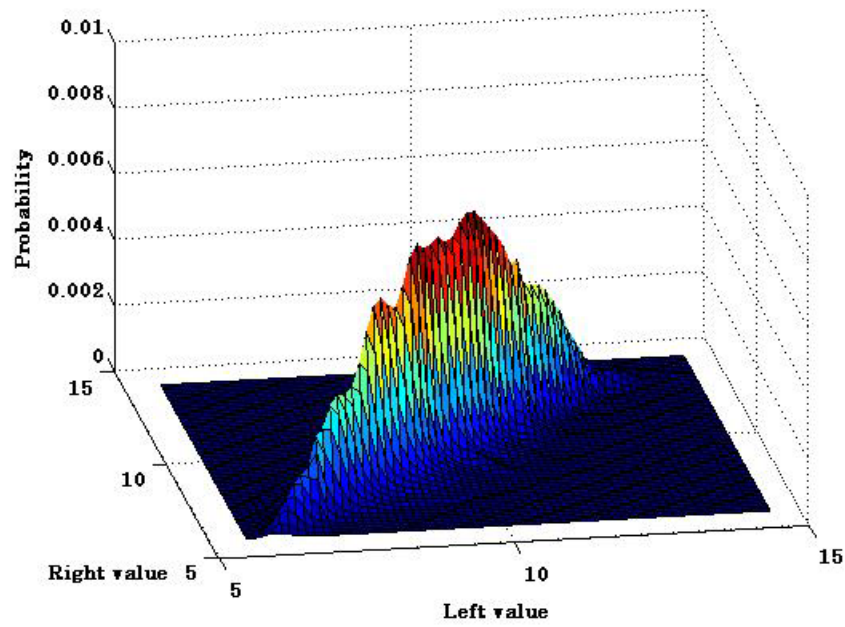
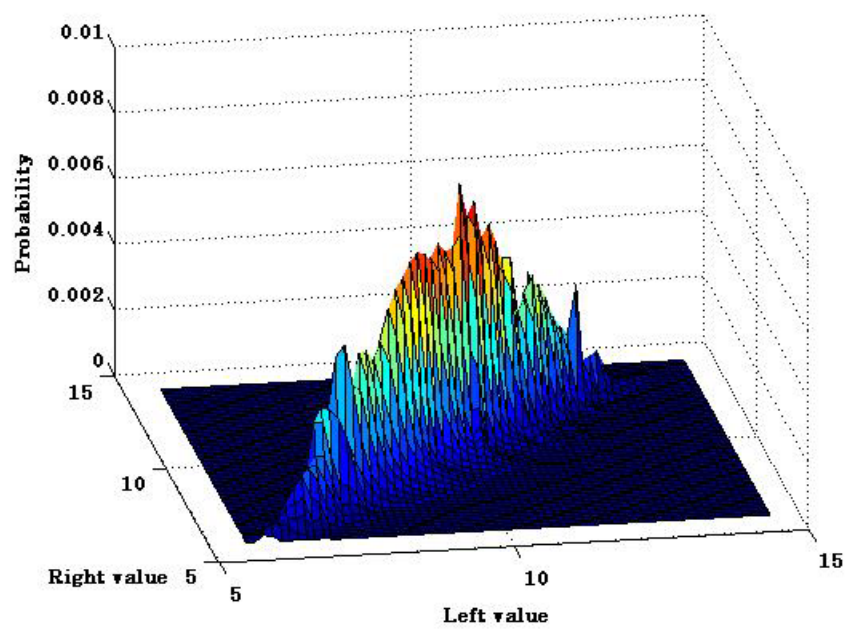


Fig. 7. (b) True value: 10



True value: 30



True value: 60



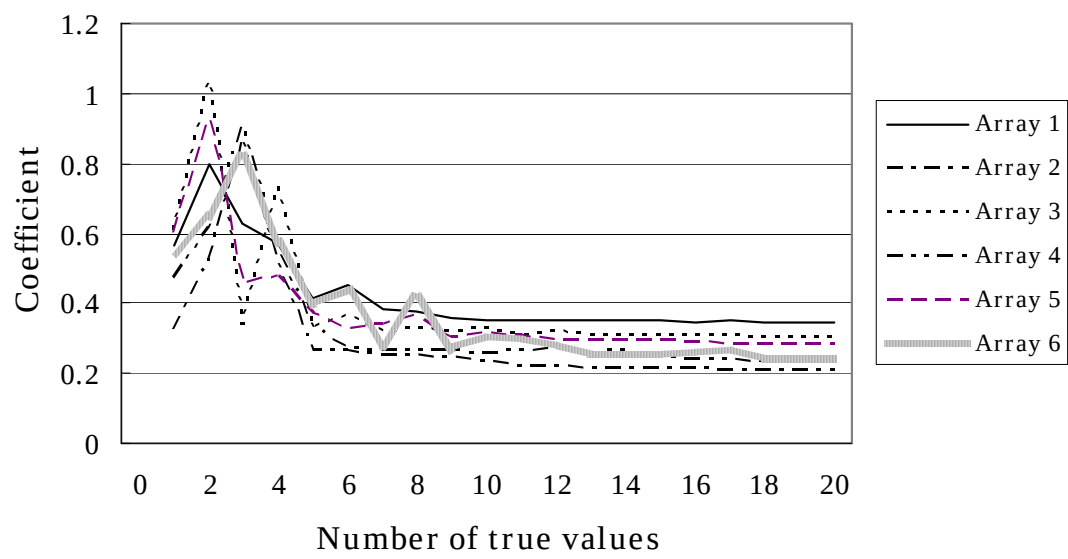


Fig. 8.

M. Takeya et al. Breadth: 8.6 cm.

Normalized true value

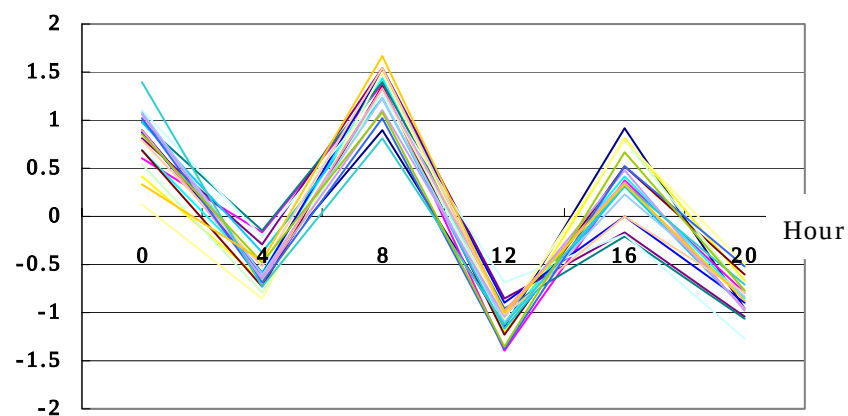
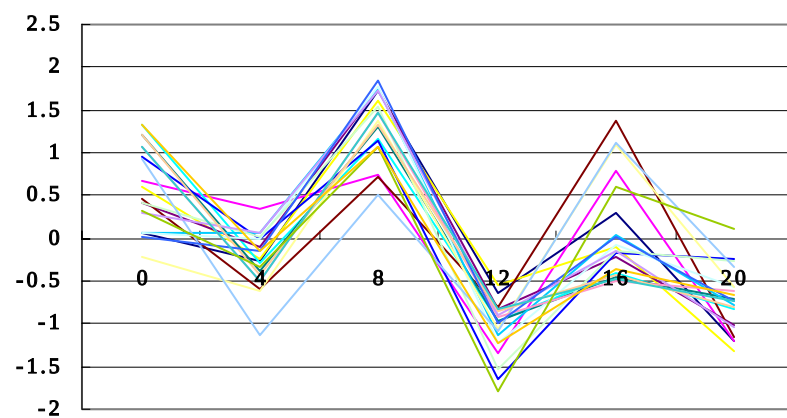
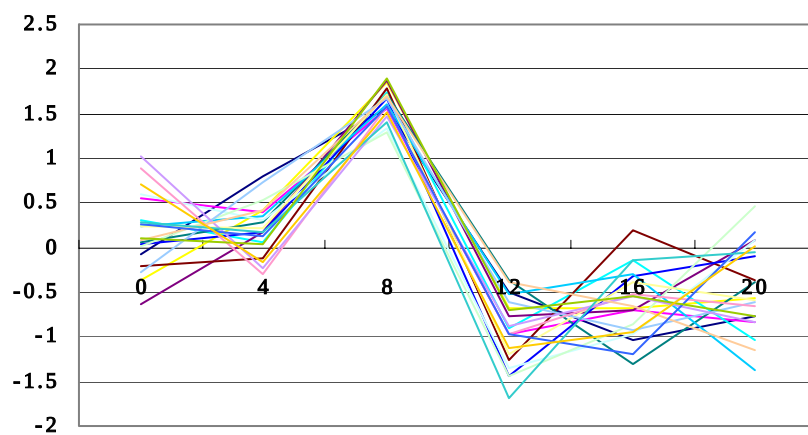


Fig. 9. Cluster 1

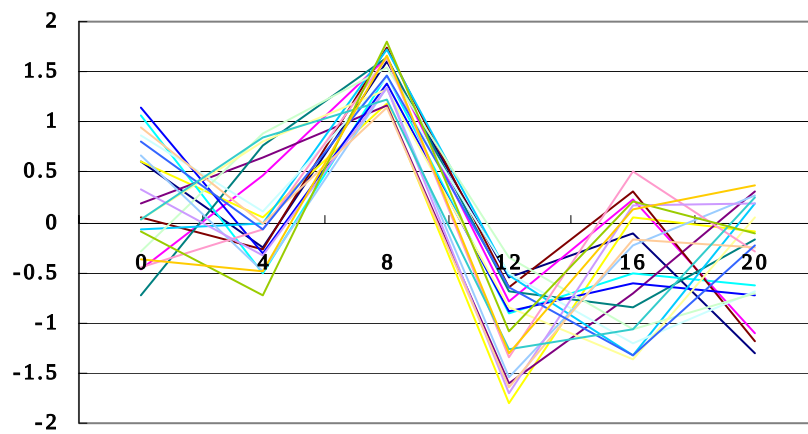
M. Takeya et al. Breadth: 4.3 cm.



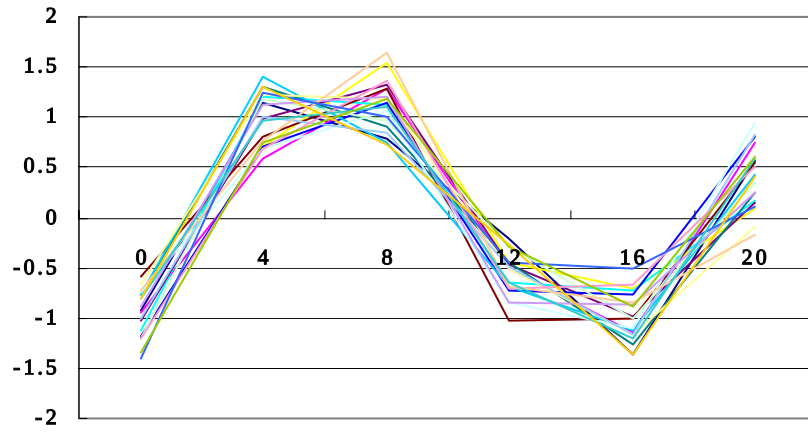
Genes having possible ranges overlapping cluster 1.



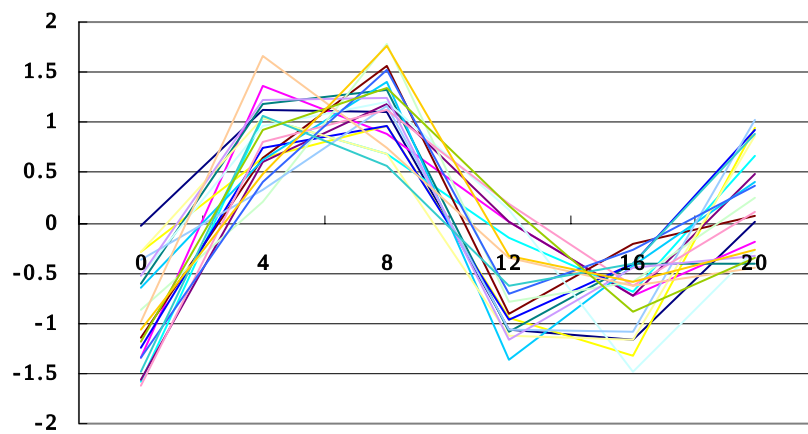
Cluster 2



Genes having possible ranges overlapping cluster 2.



Cluster 3



Genes having possible ranges overlapping cluster 3.