***List of Galaxy Jobs***

History name: ***Overlapping regions (295 Decipher patients, 19.06.14)***

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| **Job name** | **Type of CNV** | **Date** | **Notes/problems** |
| 1: DELETIONS (Decipher patients, 295, 17.06.14).txt | Deletions | 13.06.14 | OK |
| 2: DELETIONS - Sort on data 1 (on chr) |
| 3: DELETIONS - Genome Coverage BedGraph (data 1) |
| 4: DELETIONS - Join on data 1 and 3 |
| 5: DELETIONS - Group on data 4 |
| 6: DELETIONS - Join two datasets on data 5 and 3 |
| 9: DELETIONS - Cut on data 6 |
| 10: DELETIONS - Sort on data 9 (on num. of overlaps) |
| 11: DUPLICATIONS (Decipher patients, 295, 18.06.14).txt | Duplications | 13.06.14 | OK |
| 12: DUPLICATIONS - Sort on data 11 (on chr) |
| 13: DUPLICATIONS - Genome Coverage BedGraph (data 11) |
| 14: DUPLICATIONS - Join on data 11 and 13 |
| 15: DUPLICATIONS - Group on data 14 |
| 16: DUPLICATIONS - Join two Datasets on data 15 and 13 |
| 17: DUPLICATIONS - Cut on data 16 |
| 18: DUPLICATIONS - Sort on data 17 |