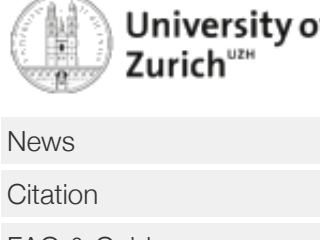


Visualization of genomic array data sets

Search Public Series	SERIES	<input type="text" value="GSE21530"/>	[?]
Search Public Arrays	ARRAY ID	<input type="text" value="GSM537760::GSM537762::GSM537765::GSM537763::GSM537767::GSM537766"/>	[?]
Plot Array Data	<input type="button" value="Show original plots & Load defaults (if existing)"/>		
Progenetix	CASE DATA PLOTTING	<input type="text" value="plot raw data values & segments"/>	[?]
 University of Zurich	GOLDEN PATH EDITION	<input type="text" value="HG18 / 36"/>	[?]
	PLOT REGION	<input type="text" value="chr5:1100000-1400000"/>	[?]
News	CHROMOSOMES TO PLOT	<input type="text" value="1:22"/>	[?]
Citation	LOSS/GAIN THRESHOLDS	<input type="text" value="-0.15"/> - <input type="text" value="0.15"/> Y AXIS FACTOR <input type="text" value="1.2"/>	[?]
FAQ & Guide	REGION SIZE	<input type="text" value="0"/> - <input type="text" value="250000"/> MIN. PROBES <input type="text" value="2"/>	[?]
Contributing Authors	EMAIL / PASSWORD	<input type="text" value="your registered email"/> <input type="text" value="your received password"/>	[?]
CompBio Zurich	LICENSE AGREEMENT	<input checked="" type="checkbox"/> I agree to license, disclaimer & citation note	
Atlas of Cytogenetics in Hematology/Oncology	<input type="button" value="Filter / replot array data"/>		

8 will be processed.

GSM537760	26 segments
GSM537761	7 segments
GSM537762	4 segments
GSM537763	1 segments
GSM537764	86 segments
GSM537765	63 segments
GSM537766	8 segments
GSM537767	50 segments

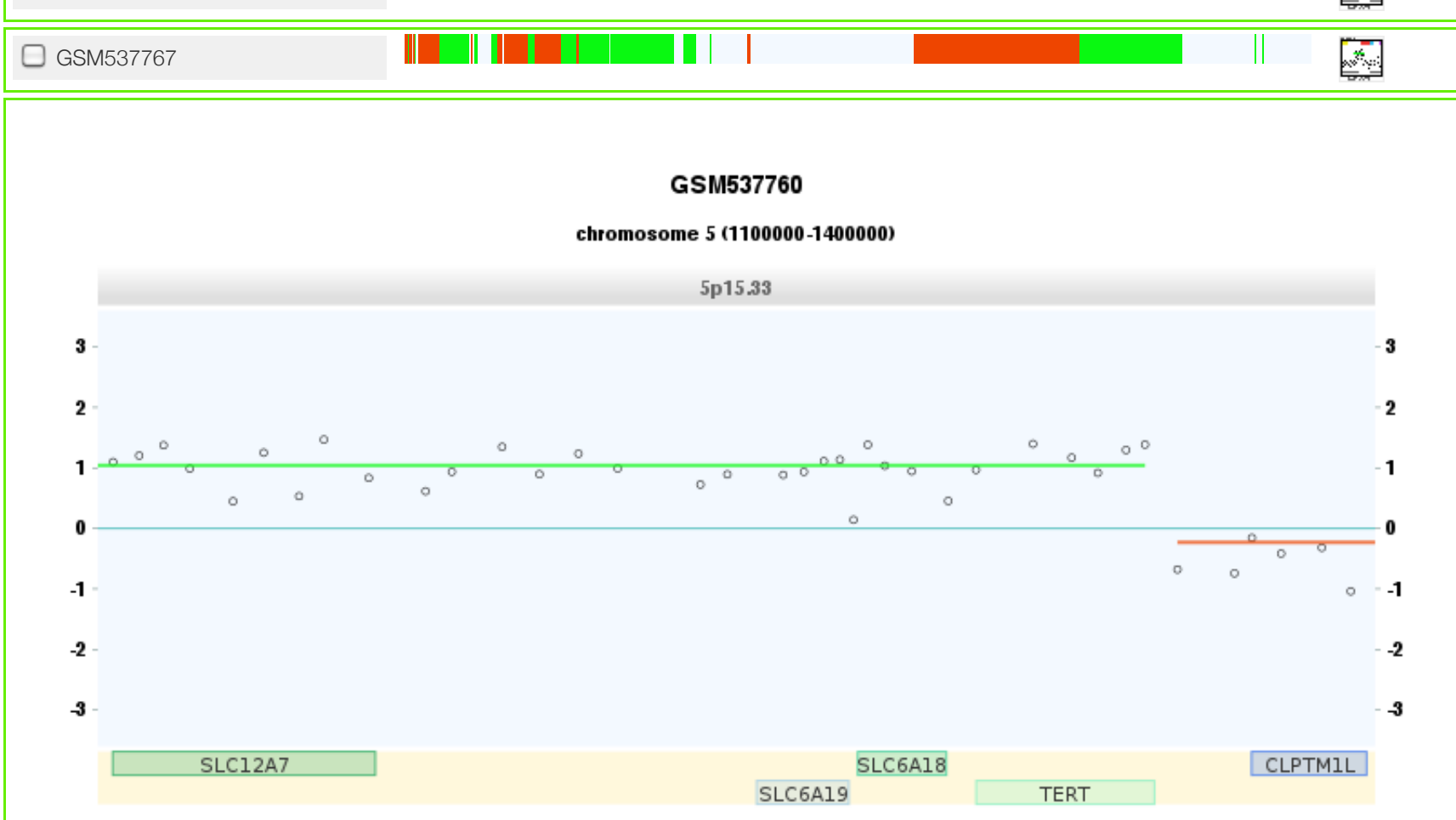
- | | | | |
|-------------------------------------|-----------|-------------------------------------|--|
| <input checked="" type="checkbox"/> | GSM537760 | Replotted array GSM537760 (1 of 8). | |
| <input checked="" type="checkbox"/> | GSM537761 | Replotted array GSM537761 (2 of 8). | |
| <input checked="" type="checkbox"/> | GSM537762 | Replotted array GSM537762 (3 of 8). | |
| <input checked="" type="checkbox"/> | GSM537763 | Replotted array GSM537763 (4 of 8). | |
| <input checked="" type="checkbox"/> | GSM537764 | Replotted array GSM537764 (5 of 8). | |
| <input checked="" type="checkbox"/> | GSM537765 | Replotted array GSM537765 (6 of 8). | |
| <input checked="" type="checkbox"/> | GSM537766 | Replotted array GSM537766 (7 of 8). | |
| <input checked="" type="checkbox"/> | GSM537767 | Replotted array GSM537767 (8 of 8). | |

REPLACE DEFAULT PLOTS save plots save defaults [?]

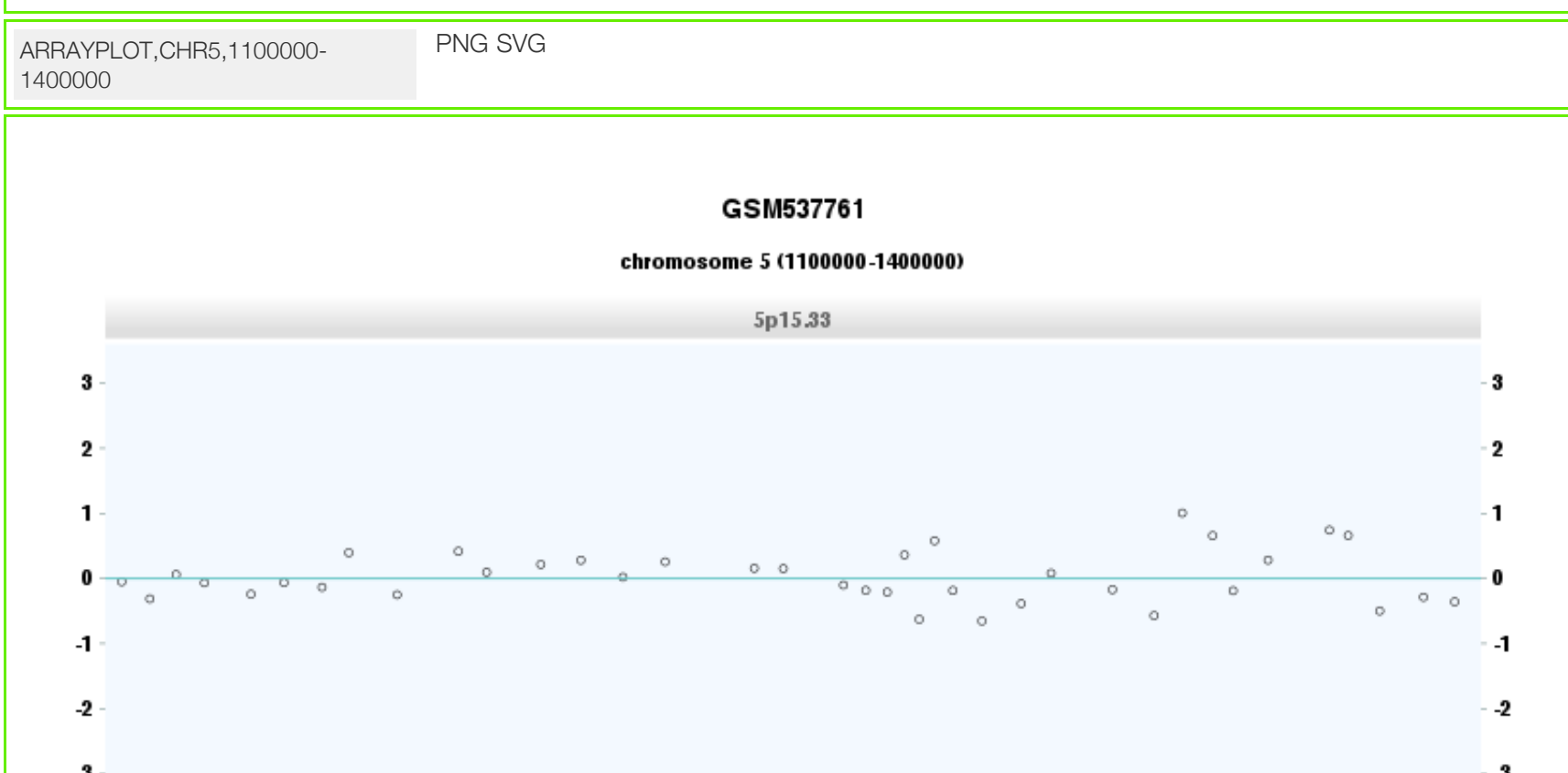
8 arrays were found.

parsing database ...

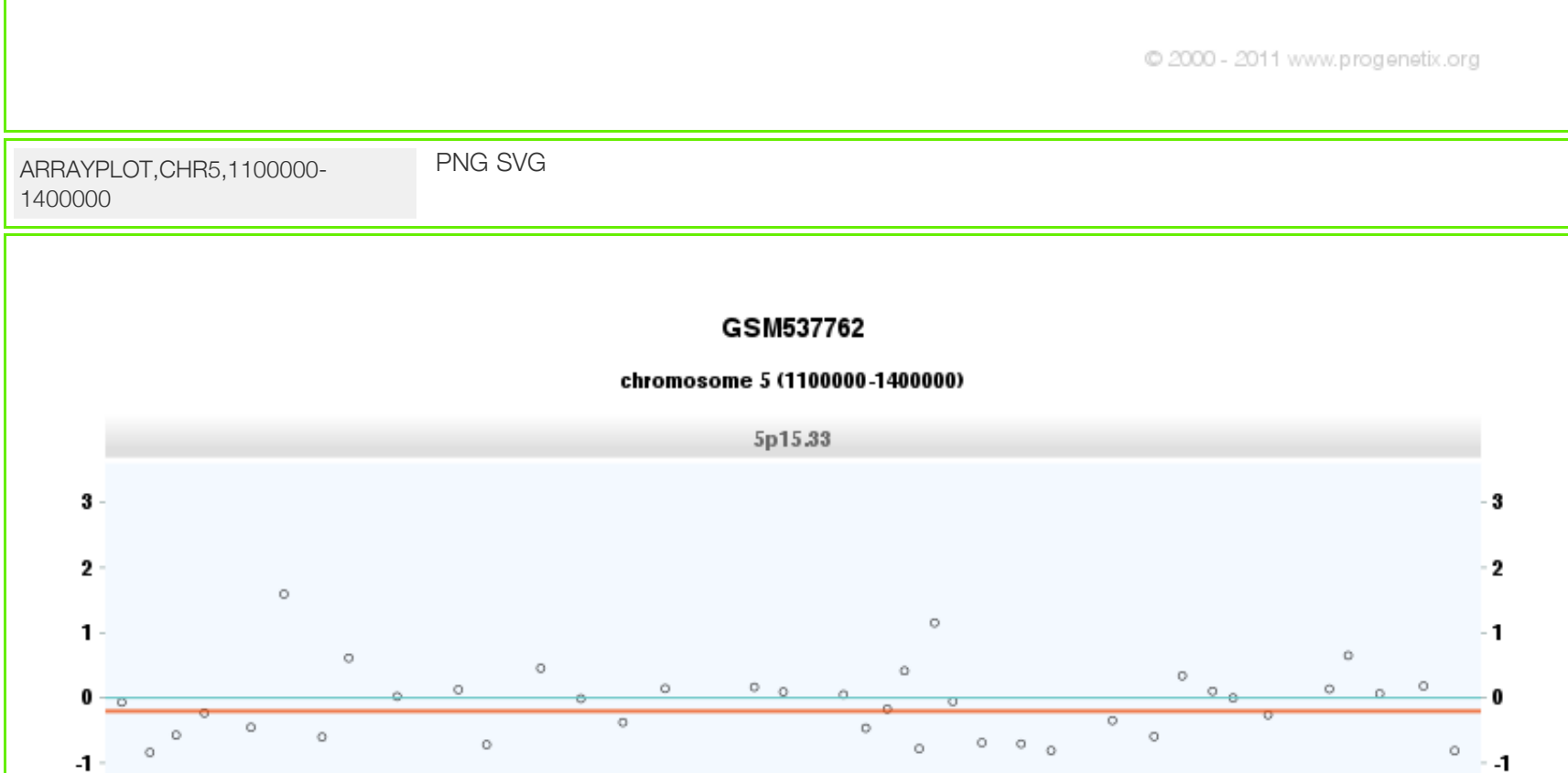
<input type="checkbox"/>	GSM537760		
<input type="checkbox"/>	GSM537761		
<input type="checkbox"/>	GSM537762		
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<input type="checkbox"/>	GSM537765		
<input type="checkbox"/>	GSM537766		
<input type="checkbox"/>	GSM537767		



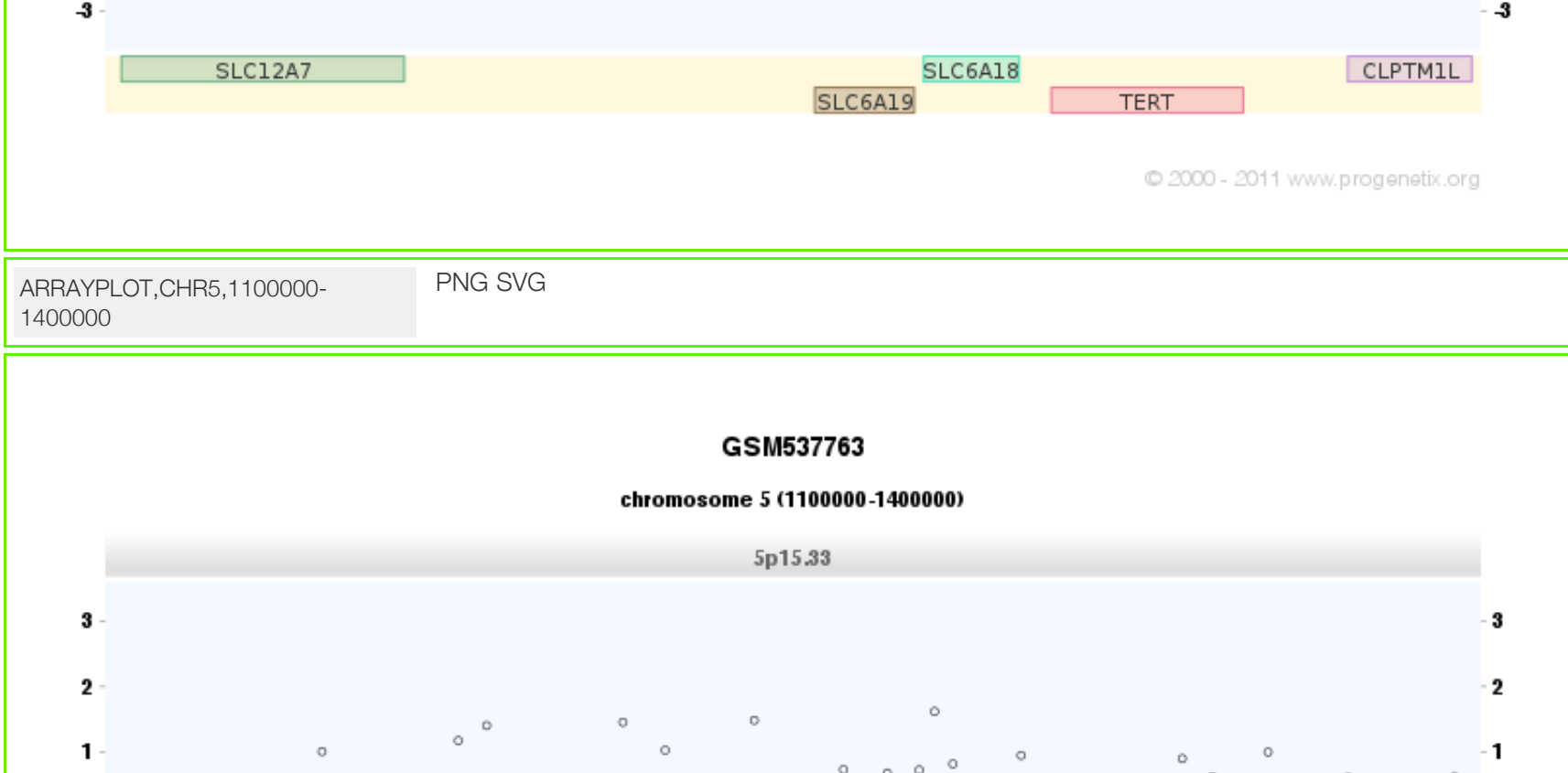
ARRAYPLOT,CHR5,1100000-1400000 PNG SVG



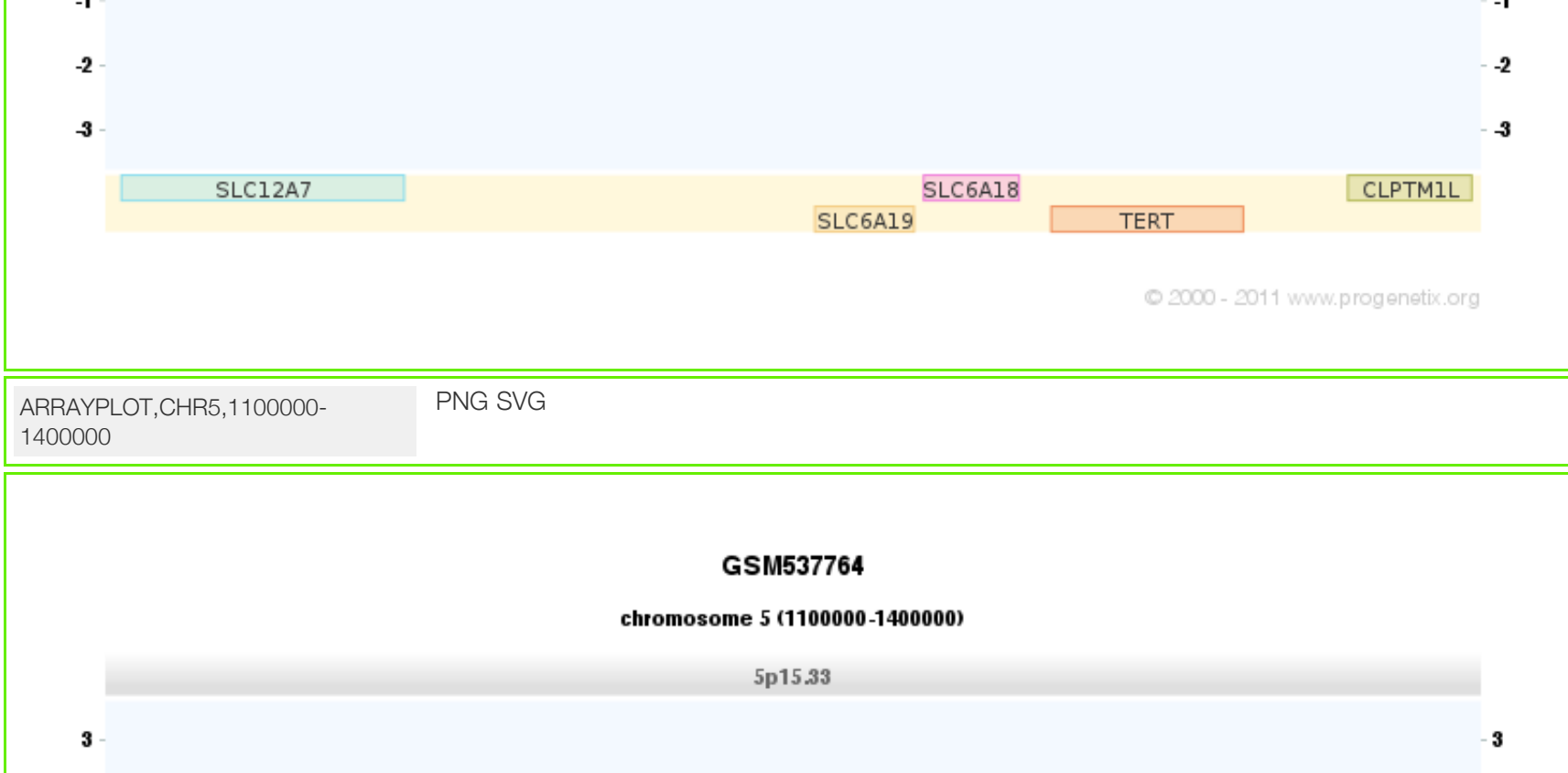
ARRAYPLOT,CHR5,1100000-1400000 PNG SVG



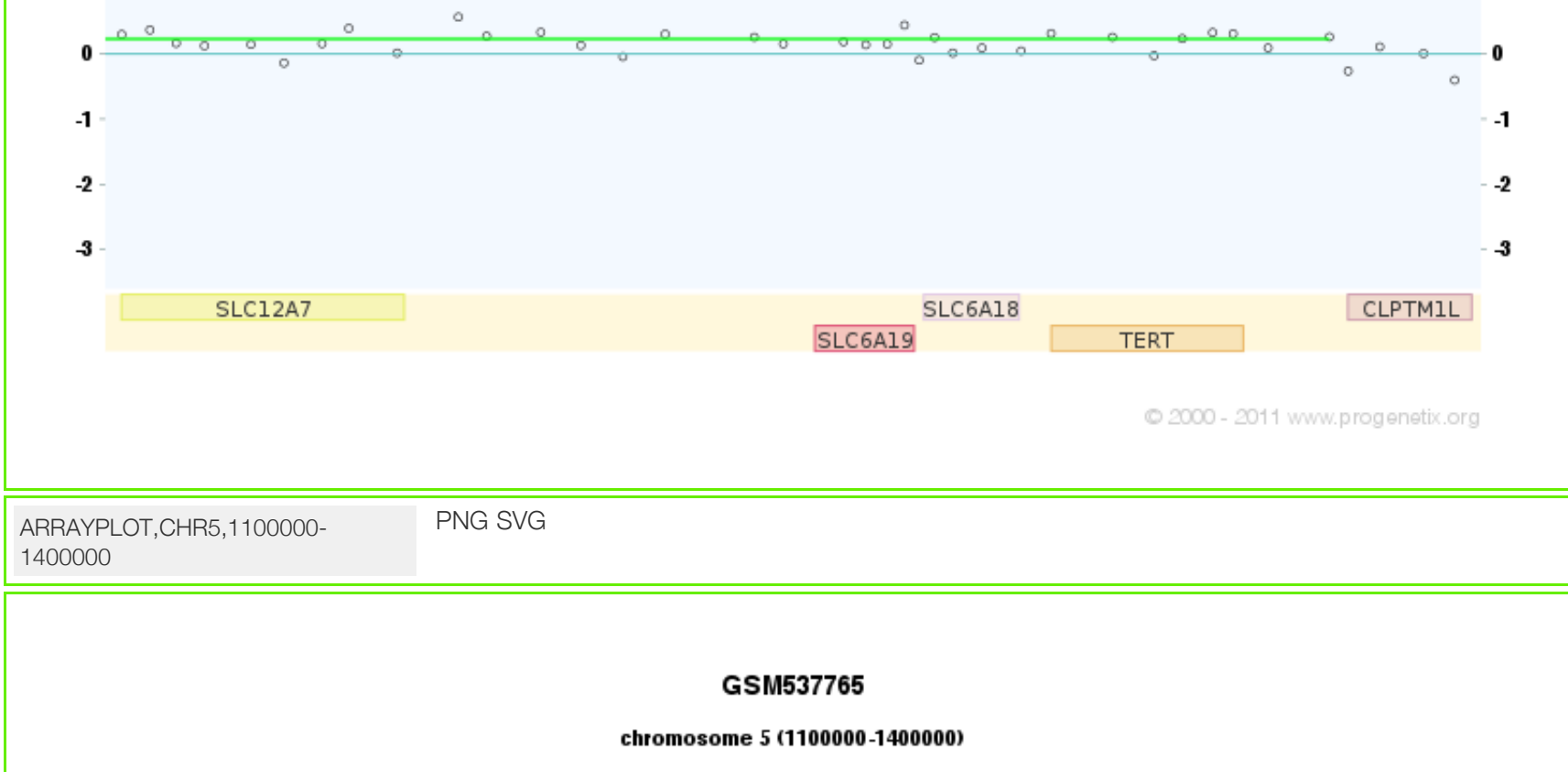
ARRAYPLOT,CHR5,1100000-1400000 PNG SVG



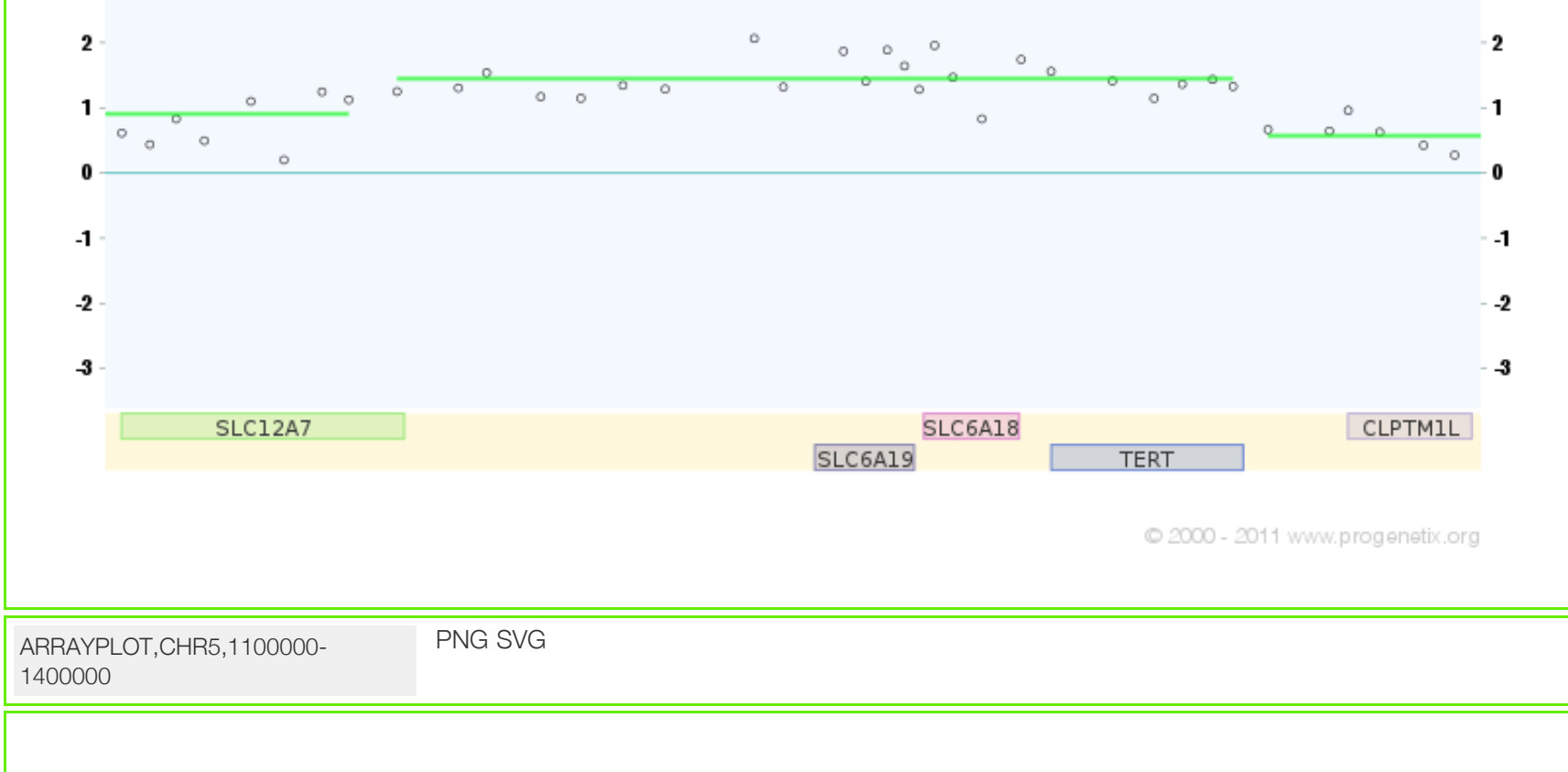
ARRAYPLOT,CHR5,1100000-1400000 PNG SVG



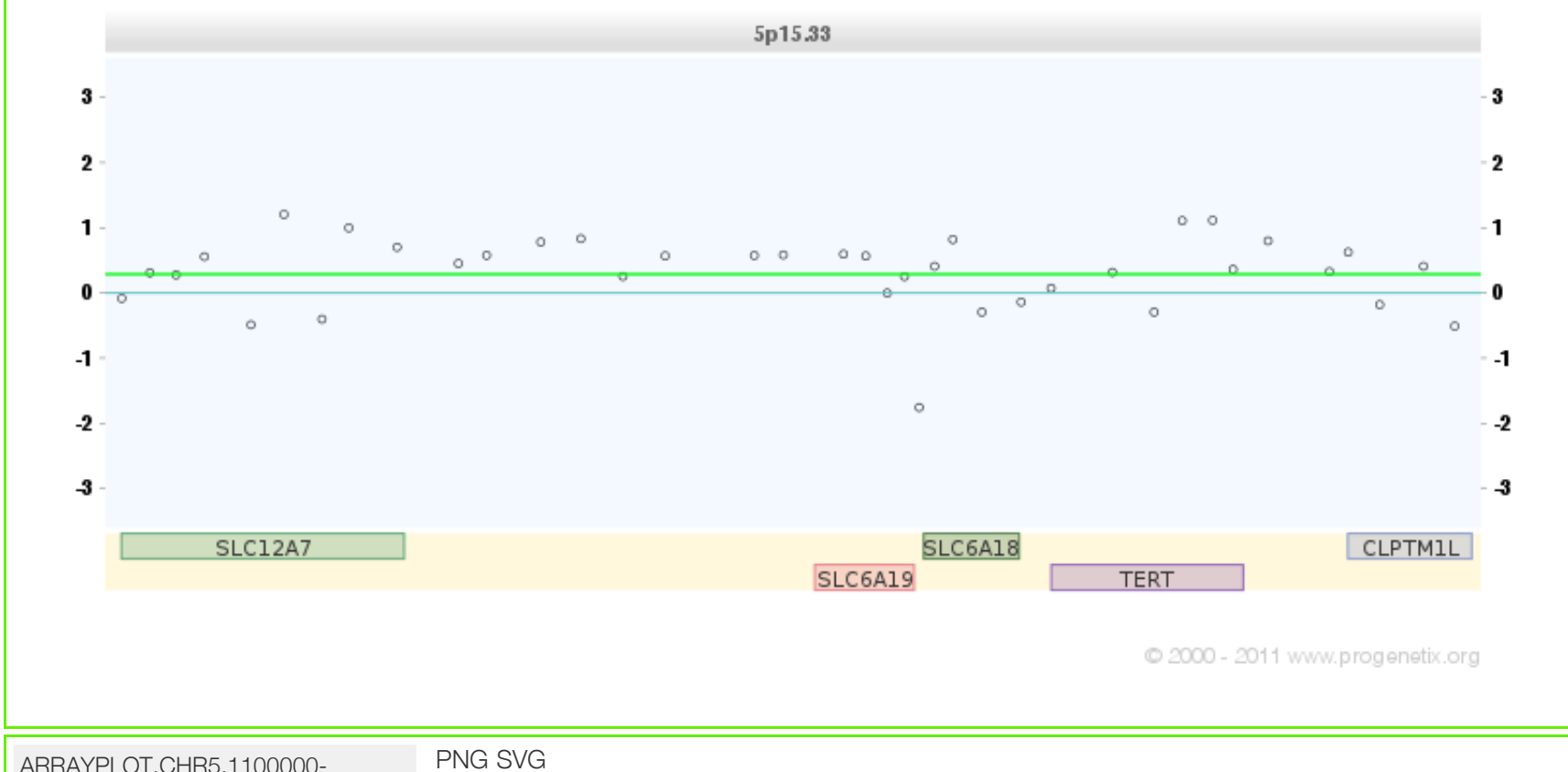
ARRAYPLOT,CHR5,1100000-1400000 PNG SVG



ARRAYPLOT,CHR5,1100000-1400000 PNG SVG



ARRAYPLOT,CHR5,1100000-1400000 PNG SVG



ARRAYPLOT,CHR5,1100000-1400000 PNG SVG

SELECT ALL [?]

These tools were developed for our research projects. You are welcome to try them out, but there is only sparse documentation. If more support and/or custom analysis is needed, please contact Michael Baudis regarding a collaborative project.