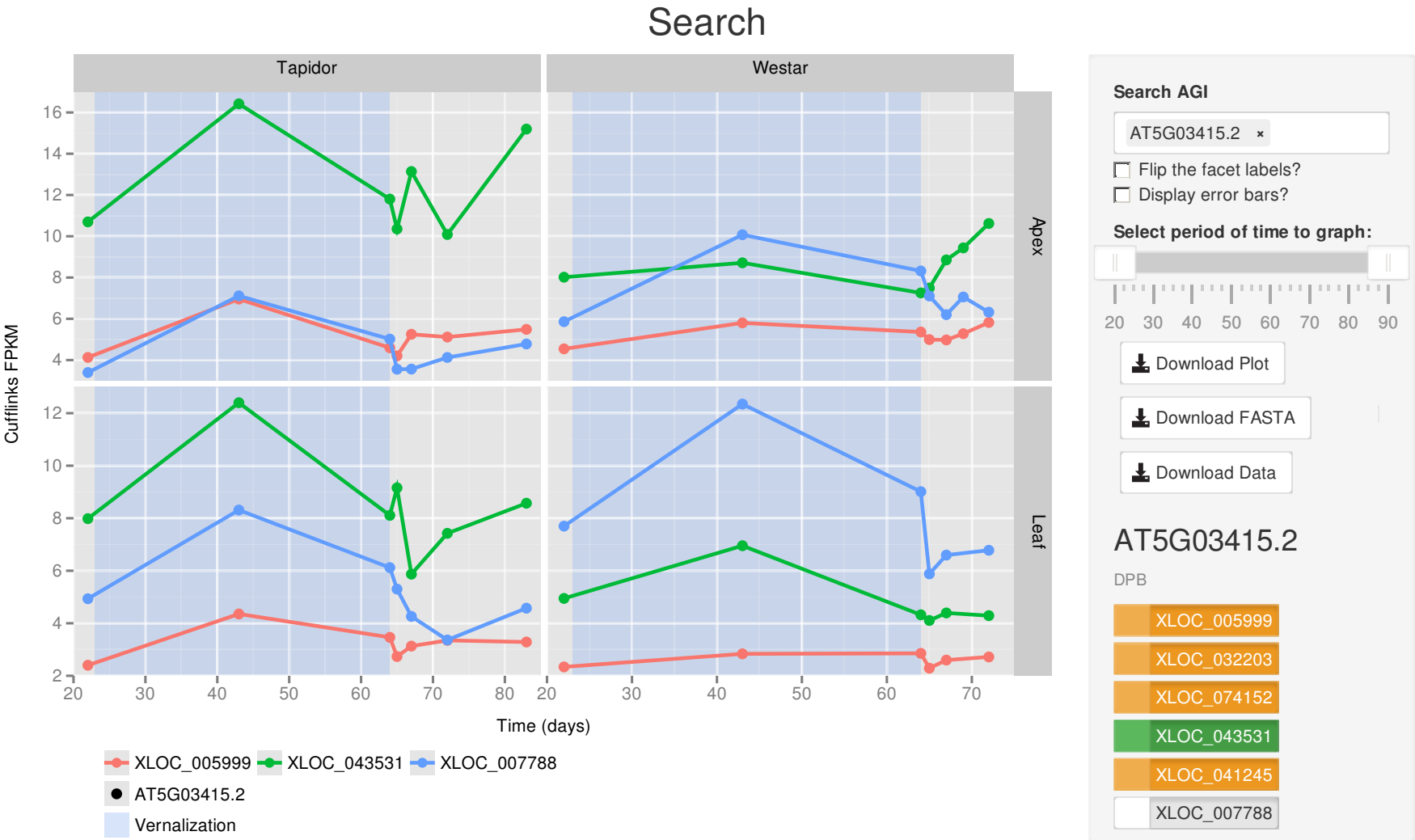


A



B

BLAST Search

Sequence:

ATGTCTATAAATATAAGAGACCCTCTTATAGTAAGCAGAGTTGTTGGAGACGTTCTTGATCCGTTTAATAGATCA
ATCACTCTAAAGGTTACTTATGGCCAAAGAGAGGTGACTAATGGCTTGGATCTAAGGCCTTCTCAGGTTCAA
AAGCCAAGAGTTGAGATTGGTGGAGAAGACCTCAGGAACCTTCTATACTTTGGTTATGGTGGATCCAGATGTTCC
AAGTCCTAGCAACCCTCACCTCCGAGAATATCTCCATTGGTTGGTGACTGATATCCCTGCTACAACCTGGAACAAC
CTTTGGCAATGAGATTGTGTGTTACGAAATCCAAGTCCCACTGCAGGAATTCATCGTGTCTGTTTATATTGTT
TCGACAGCTTGGCAGGCAAACAGTGTATGCACCAGGTTGGCGCCAGAACTTCAACACTCGCGAGTTTGCTGAG
ATCTACAATCTCGGCCTTCCCGTGGCCGCAGTTTTCTACAATTGTCAGAGGGAGAGTGGCTGCGGAGGAAGAA
GACTTTAG

There are 6 BLAST hits to the above sequence. Go to the Search tab to plot their temporal expression patterns.

C

Table

Showing 10 entries

Search:

	Brassica napus Gene	Chromosome	Start (bp)	End (bp)	Arabidopsis	Abbreviation	BLAST Identity	BLAST HSP Bit Score	BLAST HSP Length
+	XLOC_005999	chrA03	380874	384250	AT5G03415.2	DPB	80.2116402116	865.1	945
-	XLOC_007788	chrA03	381875	387292	AT5G03415.2	DPB	90	203.264	150
ArabidopsisAbbreviationBLAST IdentityBLAST HSP Bit ScoreBLAST HSP Length									
AT5G03430.189.48069241011995.811502									
AT5G03415.2DPB90203.264150									
AT5G03415.1DPB; ATDPB90203.264150									
-	XLOC_043531	chrC03	544942	545460	AT5G03415.2	DPB	80.122324159	289.825	327
ArabidopsisAbbreviationBLAST IdentityBLAST HSP Bit ScoreBLAST HSP Length									
AT5G03415.2DPB80.122324159289.825327									
AT5G03415.1DPB; ATDPB80.122324159289.825327									