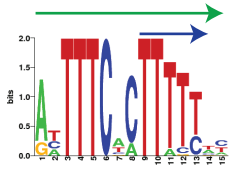
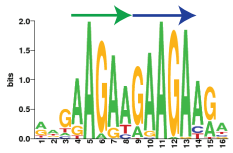
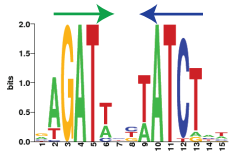
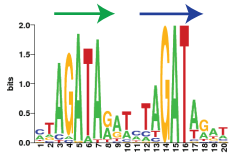
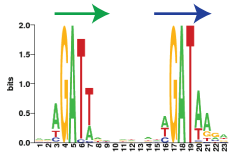
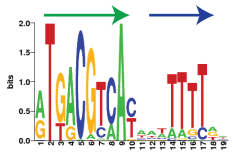
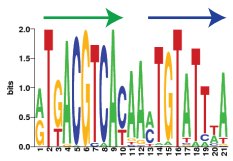
Composite Motif (CM)	ChIP TF →	Partner TF(s) →	CM Dataset Enrichment (log_2FoldChange)	Comments
	BLMP-1 (C2H2 ZF)	MAB-23 (DM) POP-1(Sox)	MAB-23 OLAP 5 1F2F (EM:3.56) MAB-23 OLAP 5 1F2F (L1:2.98) POP-1 OLAP 5 1F2F (L1:3.32) POP-1 OLAP 5 1F2F (EM:3.83)	Potentially competitive binding sites
	CEH-26 (Prospero)	CEH-26 (Prospero)	CEH-26 OLAP 4 1F2F (LE:1.89)	
	ELT-1 (GATA)	EGL-43 (C2H2 ZF) END-3 (GATA)	EGL-43 OLAP 5 1F2R (L3:2.86) END-3 OLAP 4 1F2R (L3:3.32)	
	ELT-1 (GATA)	END-1 (GATA) END-3 (GATA)	END-1 GAP 0 1F2F (L3:2.07) END-3 GAP 1 2F1F (L2:2.68)	Tandem GATA sites similar to *ges-1* promote region (Hawkins & McGhee, 1995)
	ELT-1 (GATA)	END-1 (GATA)	END-1 GAP 3 1F2F (L3:2.76) END-1 GAP 4 2F1F (L3:2.57)	Tandem GATA sites similar to *ges-1* promote region (Hawkins & McGhee, 1995)
	F23F12.9 (bZIP)	C07E3.6 (Homeodomain) CEH-51 (Homeodomain) CEH-52 (Homeodomain) MAB-23 (DM) POP-1(Sox)	C07E3.6 OLAP 3 2F1F (EM:3.88) CEH-51 OLAP 3 1F2F (EM:3.28) CEH-52 OLAP 2 1F2F (EM:3.35) MAB-23 OLAP 2 2F1F (EM:3.57) MAB-23 GAP 1 1F2F (EM:2.61) MAB-23 GAP 0 1F2F (EM:3.48) POP-1 GAP 2 1F2F (EM:3.12)	
	F23F12.9 (bZIP)	C07E3.6 (Homeodomain) CEH-26 (Prospero) CEH-34 (Homeodomain) CEH-36 (Homeodomain) CEH-51 (Homeodomain) CEH-52 (Homeodomain) DMD-3 (DM) DMD-4 (DM) DMD-5 (DM) DMD-6 (DM) FKH-9 (Forkhead) GEI-3 (Sox) MAB-23 (DM) NHR-168 (NHR) NHR-42 (NHR) NHR-66 (NHR) SCRT-1 (C2H2 ZF) SYD-9 (C2H2 ZF)	C07E3.6 GAP 3 2F1F (EM:4.22) CEH-26 GAP 2 1F2F (EM:4.53) CEH-34 GAP 1 2F1F (EM:4.8) CEH-36 GAP 3 1F2F (EM:5.15) CEH-51 GAP 3 1F2F (EM:3.62) CEH-52 GAP 2 2F1F (EM:3.22) DMD-3 GAP 1 2F1F (EM:4.65) DMD-4 GAP 2 2F1F (EM:5.62) DMD-5 OLAP 4 1F2F (EM:4.64) DMD-5 GAP 1 2F1F (EM:5.18) DMD-6 GAP 0 2F1F (EM:3.58) FKH-9 GAP 2 1F2F (EM:4.8) GEI-3 GAP 2 2F1F (EM:4.37) MAB-23 GAP 4 1F2F (EM:3.18) NHR-168 GAP 0 2F1F (EM:5.38) NHR-42 GAP 1 2F1F (EM:4.72) NHR-66 OLAP 1 2F1F (EM:5.93) SCRT-1 GAP 1 1F2F (EM:3.24) SCRT-1 GAP 3 2F1F (EM:3.32) SCRT-1 GAP 4 2F1F (EM:3.23) SYD-9 GAP 0 1F2F (EM:4.75)	