## Genes gained in Stylomatophora BP TreeMap

positive regulatior of primary cell septum biogenesi	of septa	tion ch	gulation of romosome ganization	positive regulation of mitotic nuclear division	ascospore formation	on	r	cytoplasmic microtubule organization		oocyte karyosome formation		totic spindle rganization	heterochromatin assembly
mitotic metaphase plate congression	regulation of mitotic nuclear division	regulation of mitotic spindle organization	attachment mitotic spind microtubules kinetochor	dle regulation of mitotic cytokinetic	synaptonemal		fusom	e organization <mark>meiotic ce</mark>	meiosis I II cycle		osome gation	male meiosis I	positive regulation of DNA methylation–dependent heterochromatin assembly
metaphase/anaphase transition of cell cycle	l <b>regulation of sis</b> metaphase/anaphase transition of cell cycle	positive regulation ster chromatid s metaphase/anaphase transition	positive egregation mitotic siste chromatid separation	segregation	complex assembl	У	meiotic cell cycle		microtubule cytoskeletor organization	toskeleton cell cy		meiotic chromosome segregation	regulation of DNA methylation–dependent heterochromatin assembly
metaphase/anaphase transition of mitotic cell cycle	positive regulation of mitotic cytokinesis	modulation l symbiont o host cell cyc	by modulation f virus of h	n by regulation of ost chromosome	cell division		fema	ale meiosis I	reciprocal meio recombinatio		ule microt	indle nuclear ubules chromosom segregation	
positive regulation of chromatin organization	positive regulation of chromosome separation	regulation cytokinetic		regulation of heterochromatin organization	piRNA metabolic	DNA	JA	reciprocal homol recombination	on reco	nologous mbination	double-str break repair v	recombinationa ia repair	DNA methylation–dependent heterochromatin assembly
positive regulation of heterochromatin organization	modification by virus of host ce cycle regulation	ll cellular co	mponent	regulation of chromatin organization	process	recombina	DNA recom		nd regula	regulation of DNA recombination		uble-strand eak repair	regulation of heterochromatin assembly