Comp	name	usage	icon
geom	geom_rect	rectangle	figure/geom_rect.pdf
	$geom\_segment$	segment	figure/geom_segment.pdf
	geom_chevron	chevron	figure/geom_chevron.pdf
	$geom\_arrow$	arrow	figure/geom_arrow.pdf
	$geom\_arch$	arches	figure/geom_arch.pdf
	geom_bar	bar	figure/geom_bar.pdf
	$geom\_alignmer$	ntgene structure	figure/geom_alignment.pdf
stat	stat_aggregate	aggregation in sliding window	figure/stat_aggregate.pdf
	$stat\_coverage$	coverage for reads	figure/stat_coverage_icon.pdf
	$stat\_gene$	gene structure	figure/stat_gene.pdf
	stat_identity	identity mode	figure/stat_identity.pdf
	$stat\_mismatch$	mismatch summary for variants	figure/stat_mismatch.pdf
	$stat\_stepping$	stepping levels	figure/stat_stepping.pdf
	stat_table	tabulate ranges	figure/stat_table.pdf
coord	linear	default linear but facet by chromosomes space	figure/coord_linear.pdf
	genome	put everything on genome coordiante	figure/coord_genome.pdf
	truncate	compact view by cutting	figure/coord_truncate_gaps.pdf
	gaps	gaps	
layout	default	normal layout	figure/coord_linear.pdf
	karyogram	karyogram layout	figure/layout_karyogram.pdf
	circle	circular layout	figure/layout_circle.pdf
faceting	g formula	formula based faceting	figure/facet.pdf
	ranges	ranges based faceting	figure/facet_gr.pdf
scale	not extended	ggplot2 default	

Table 1: Components of the basic grammar of graphics, with the extensions available in ggbio.

object	usage	icon
GRanges	Genomic intervals	figure/geom_rect.pdf
GRangesList	Genomic interval	figure/grl.pdf
	list	
IRanges	intervals	figure/stat_stepping.pdf
Gapped Alignments	gapped alignments	figure/stat_stepping.pdf
BamFile	alignemnts files	figure/stat_coverage_icon.pdf
character	for flat files	figure/geom_bar.pdf
Rle/RleList	atomic vector	figure/stat_stepping.pdf
TranscriptDb	gene structure	figure/stat gene.pdf
VCF	Variant calling for-	figure/vcf.pdf
	mat	

Table 2: Formal data model and their default graphics