

ENCODE nets Review summary

	Section	# of words	% of words	# negative comments	% of negative comments	# positive comments	% of positive comments	# of sub figures	% of figures
1	Abstract	249	2.78%	1	1.85%	0	0.00%	0	0.00%
2	Introduction	759	8.49%	1	1.85%	0	0.00%	0	0.00%
3	Overview of data and data processing	543	6.07%	1	1.85%	0	0.00%	0	0.00%
4	The combinatorial nature of TF co-association	802	8.97%	2	3.70%	0	0.00%	3	14.29%
5	Co-association within the focus-factor domain	1061	11.86%	6	11.11%	1	12.50%	3	14.29%
6	Summarizing co-association of transcription factors	561	6.27%	3	5.56%	0	0.00%	1	4.76%
7	Assembly of pairwise interactions into hierarchies	530	5.93%	12	22.22%	0	0.00%	3	14.29%
8	Connecting the promoter-regulation hierarchy to other networks to create a complex meta-network	670	7.49%	1	1.85%	1	12.50%	1	4.76%
9	Relating TF network connectivity to additional genomic properties	1107	12.38%	14	25.93%	0	0.00%	1	4.76%
10	Collaboration between levels of the regulatory hierarchy	380	4.25%	4	7.41%	0	0.00%	2	9.52%
11	The regulatory network contains many enriched motifs	620	6.93%	2	3.70%	1	12.50%	4	19.05%
12	Analyzing allele-specific binding in a network context	544	6.08%	0	0.00%	0	0.00%	2	9.52%
13	Analyzing selection in a network context	414	4.63%	0	0.00%	0	0.00%	1	4.76%
14	Discussion	705	7.88%	0	0.00%	0	0.00%	0	0.00%
15	General		0.00%	5	9.26%	5	62.50%	0	0.00%
16	Supplamentery		0.00%	2	3.70%	0	0.00%	0	0.00%
	total	8945	100.00%	54	100.00%	8	100.00%	21	100%

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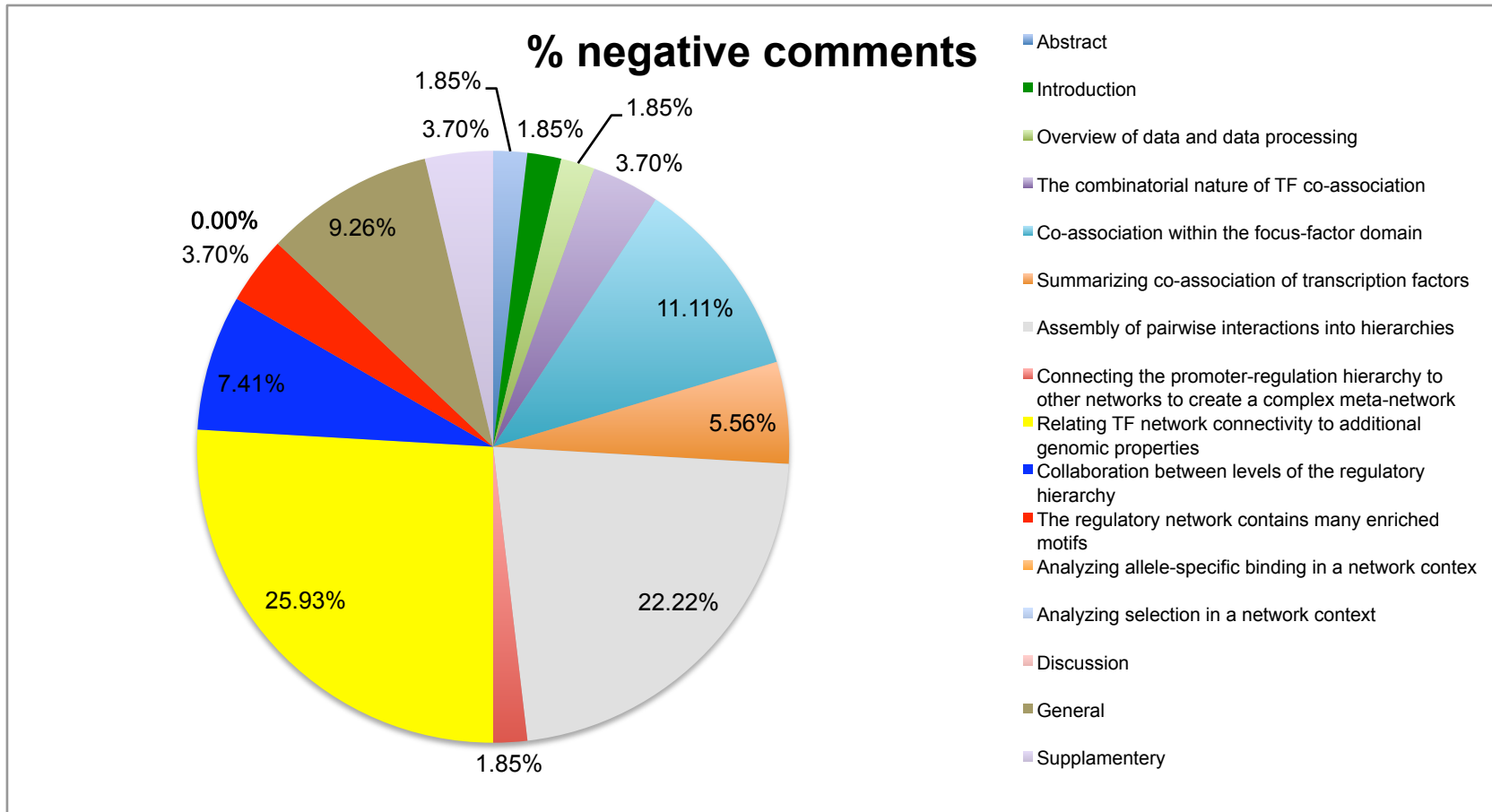
Figures

Fig	Section number
1A	The combinatorial nature of TF co-association
1B	The combinatorial nature of TF co-association
1C	The combinatorial nature of TF co-association
1D	Co-association within the focus-factor domain
1E	Co-association within the focus-factor domain
2A	Summarizing co-association of transcription factors
2B	Co-association within the focus-factor domain
3A	Assembly of pairwise interactions into hierarchies
3B	Assembly of pairwise interactions into hierarchies
3C	Assembly of pairwise interactions into hierarchies
3D	Connecting the promoter-regulation hierarchy to other networks to create a complex meta-network
4A	Relating TF network connectivity to additional genomic properties
4B	Collaboration between levels of the regulatory hierarchy
4C	Collaboration between levels of the regulatory hierarchy

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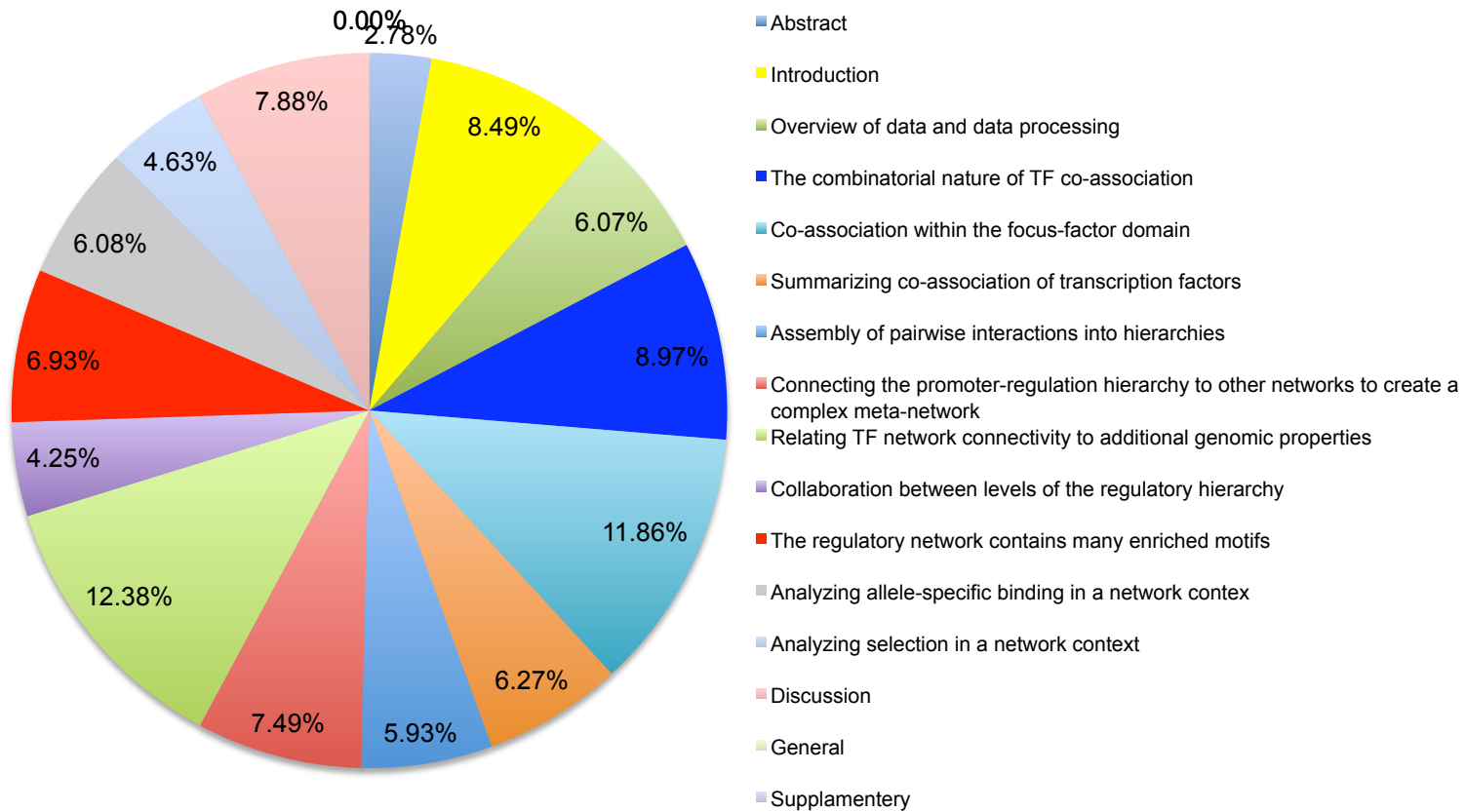
5A	The regulatory network contains many enriched motifs
5B	The regulatory network contains many enriched motifs
5C	The regulatory network contains many enriched motifs
5D	The regulatory network contains many enriched motifs
6A	Analyzing allele-specific binding in a network context
6B	Analyzing allele-specific binding in a network context
6C	Analyzing selection in a network context

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% of words in each section



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