stelpflug2016 B73 [93]			654		
walley2016 B73 [23]			514		
briggs B73 [23]			441		
briggs Mo17 [23]			547		
briggs BxM [23]	-		691		
leaf dev leaf [40]			798		
seed dev embryo_endpsperm [61]	-		1205		
dev atlas combined [139]			521		
root dev root [50]	-	_	4474		
eitchen2013 seedling_leaf3 [62]	-		420		
hirsch2014 seedling [503]			254		
leiboff2015 SAM [383]			359		
jin2016 kernel [368]			101		
lin2017 ear [26]	-		547		
lin2017 root [27]			508		$\overline{}$
lin2017 shoot [27]			614		
lin2017 tassel [26]			768		$\overline{}$
lin2017 SAM [27]			628		
kremling2018 GRoot [274]	-		309		
kremling2018 GShoot [280]			444		
kremling2018 Kern [231]			465		
kremling2018 L3Base [264]			140		
kremling2018 L3Tip [266]			358		
kremling2018 LMAD [204]	-		374		
kremling2018 LMAN [262]	-		219		
shaefer2018 root_GCN [48]			447		
kaeppler2018 endosperm [89]			2044		
kaeppler2018 internode [43]			1609		
kaeppler2018 leaf [50]			1745		
kaeppler2018 root [49]			1936		
kaeppler2018 seedling [216]	-		556		
kaeppler2018 seedling inbred [166]	-		487		
kaeppler2018 seedling hybrid [50]			2141		
biomap endosperm [121]			1734		
biomap internode [77]			1299		
biomap leaf [84]			1829		
biomap root [84]			1269		
biomap seedling [84]			1579		
lin2017 5*27 [133]			693		
kremling2018 7*306 [1781]			304		
kaeppler2018 5*96 [447]			1030		
briggs 3*23 [73]			10246		$\overline{}$
biomap 5*121 [450]			1225		= =
li2013 SAM [107]			2105		
walley2016 mRNA [23]			510		
huang2018 leaf [394]			743		
huang2018 root [176]			1106		
huang2018 SAM [406]			483		
huang2018 seed [159]			902		
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	-1.0	-0.5	0.0	0.0	1.0