GENCODE v27	100.0% (15,778/15,778)	98.0% (15,474/15,778)	99.0% (15,684/15,778)	70.0% (11,125/15,778)	47.0% (7,489/15,778)	87.0% (13,877/15,778)	58.0% (9,168/15,778)	96.0% (15,191/15,778)	37.0% (5,902/15,778)	47.0% (7,513/15,778)	95.0% (15,028/15,778)
GENCODE v47	42.0% (15,265/35,934)	100.0% (35,934/35,934)	74.0% (26,808/35,934)	31.0% (11,324/35,934)	30.0% (11,124/35,934)	37.0% (13,353/35,934)	47.0% (16,963/35,934)	59.0% (21,229/35,934)	24.0% (8,640/35,934)	31.0% (11,183/35,934)	69.0% (24,999/35,934)
IncRNA-merge	15.0% (17,449/109,892)	29.0% (32,057/109,892)	100.0% (109,892/109,892)	15.0% (16,998/109,892)	31.0% (34,764/109,892)	15.0% (16,725/109,892)	53.0% (58,696/109,892)	67.0% (74,340/109,892)	15.0% (17,497/109,892)	14.0% (16,406/109,892)	77.0% (84,866/109,892)
bigTranscriptome	81.0% (11,355/14,005)	83.0% (11,666/14,005)	100.0% (14,005/14,005)	100.0% (14,005/14,005)	52.0% (7,344/14,005)	82.0% (11,532/14,005)	60.0% (8,482/14,005)	85.0% (11,917/14,005)	37.0% (5,224/14,005)	46.0% (6,536/14,005)	95.0% (13,336/14,005)
fantomCat	24.0% (7,219/29,587)	39.0% (11,730/29,587)	100.0% (29,587/29,587)	24.0% (7,276/29,587)	100.0% (29,587/29,587)	23.0% (7,092/29,587)	50.0% (15,087/29,587)	51.0% (15,174/29,587)	19.0% (5,772/29,587)	22.0% (6,651/29,587)	67.0% (19,975/29,587)
GENCODEv20+	95.0% (13,279/13,861)	93.0% (13,028/13,861)	100.0% (13,861/13,861)	77.0% (10,798/13,861)	49.0% (6,805/13,861)	100.0% (13,861/13,861)	57.0% (8,006/13,861)	93.0% (12,913/13,861)	38.0% (5,304/13,861)	47.0% (6,642/13,861)	93.0% (12,940/13,861)
miTranscriptome	17.0% (9,076/52,141)	33.0% (17,454/52,141)	100.0% (52,141/52,141)	16.0% (8,843/52,141)	28.0% (15,120/52,141)	16.0% (8,554/52,141)	100.0% (52,141/52,141)	41.0% (21,487/52,141)	16.0% (8,708/52,141)	18.0% (9,834/52,141)	87.0% (45,557/52,141)
NONCODE	22.0% (17,211/77,407)	35.0% (27,685/77,407)	100.0% (77,407/77,407)	18.0% (14,314/77,407)	24.0% (18,973/77,407)	20.0% (15,791/77,407)	38.0% (29,893/77,407)	100.0% (77,407/77,407)	15.0% (11,835/77,407)	17.0% (13,196/77,407)	79.0% (61,742/77,407)
refSeq	37.0% (5,754/15,431)	58.0% (8,995/15,431)	100.0% (15,431/15,431)	33.0% (5,110/15,431)	37.0% (5,765/15,431)	35.0% (5,508/15,431)	57.0% (8,868/15,431)	62.0% (9,592/15,431)	100.0% (15,431/15,431)	57.0% (8,889/15,431)	93.0% (14,376/15,431)
CHESS v3.1.2	41.0% (7,279/17,365)	66.0% (11,584/17,365)	84.0% (14,629/17,365)	36.0% (6,300/17,365)	38.0% (6,652/17,365)	39.0% (6,845/17,365)	56.0% (9,847/17,365)	63.0% (10,971/17,365)	50.0% (8,830/17,365)	100.0% (17,365/17,365)	82.0% (14,407/17,365)
LncBook v2.0	17.0% (16,813/95,415)	31.0% (30,182/95,415)	89.0% (85,578/95,415)	16.0% (15,938/95,415)	24.0% (23,108/95,415)	16.0% (15,585/95,415)	55.0% (52,841/95,415)	62.0% (59,272/95,415)	16.0% (16,131/95,415)	17.0% (16,265/95,415)	100.0% (95,415/95,415)
	GENCODE v27	GENCODE v47	IncRNA-merge	bigTranscriptome	fantomCat	GENCODEv20+	miTranscriptome	NONCODE	refSeq	CHESS v3.1.2	LncBook v2.0

% of genes

75%

50%

25%