

Table S3: Data collection and processing procedures for public chromosomal alterations datasets in BTECH.

Track Description	Data Resource	Title	Methods	PubMed ID	Analysis Procedures
aCGH for Ependymomas (68 patients)	GSE3435	Identification of gains on 1q and epidermal growth factor receptor overexpression as independent prognostic markers in intracranial ependymoma	Array-CGH	16609018	Download from the GEO site
Glial Brain Tumors_GSE1991	GSE1991	High-resolution genome-wide mapping of genetic alterations in human glial brain tumors	Array-CGH	15899798	Download from the GEO site
SNPchip_178Gliomas	Publication in 2006	High-resolution global genomic survey of 178 gliomas reveals novel regions of copy number alteration and allelic imbalances	SNP chip	17018597	Extract from publication
Neuroblastoma_LOH	Publication in 2005	High-resolution detection and mapping of genomic DNA alterations in neuroblastoma	Array-CGH	15892104	Extract from publication
Recombine_Rate (decode, Marshfield, Genethon)	UCSC, released in 2006	Calculated sex-averaged rates of recombination	Genetic maps	9718341, 8600387, 12053178	Download from the UCSC site