OncoKB, MSK's Precision Oncology Knowledge Base

Sarah P. Suehnholz¹, Moriah H. Nissan¹, Hongxin Zhang¹, Ritika Kundra¹, Calvin Lu¹, Katelyn M. Mullen¹, Katelyn M. Mullen¹, Katelyn M. Mullen¹, Katelyn M. Tran¹, Yan Li^{1,3}, Maria E. Arcila^{1,4}, Marc Ladanyi⁴, Michael F. Berger^{1,4}, Julia E. Rudolph¹, Paul Sabbatini⁵, Ryan Ptashkin^{1,4}, Ahmet Dogan⁴, Ross L. Levine^{1,5}, Jianjiong Gao^{1,6}, David B. Solit^{1,5}, Nikolaus Schultz^{1,6} and Debyani Chakravarty^{1,4}



¹Kravis Center for Molecular Oncology, Memorial Sloan Kettering Cancer Center, ²Pharmacology Department of Surgery, Memorial Sloan Kettering, ⁴Department of Pathology and Laboratory Medicine, Memorial Sloan Kettering Cancer Center, 5Department of Medicine, Memorial Sloan Kettering Cancer Center, 6Computational Oncology Service, Department of Epidemiology and Biostatistics

Comprehensive evidence-based information about cancer variants Welcome to OncoKB 03/01/2022 MSK's Precision Oncology Knowledge Base An FDA-Recognized Human Genetic Variant Database* 687 5707 110 **132** Cancer Types Alterations Level 1 4 Level 4 Level R1/R2 Level 3 2 Level 2

Clinical evidence

26 Genes

An FDA-recognized* human genetic variant database

Standard care

20 Genes

FDA recognizes Memorial Sloan-Kettering database of molecular tumor marker information

FDA-approved drugs

43 Genes

"On October 7, 2021, the Food and Drug Administration granted recognition to a partial listing of the Memorial Sloan Kettering Cancer Center's Oncology Knowledge Base (OncoKB) as the first tumor mutation database to be included in the Public Human Genetic Variant Database.

* FDA recognition of OncoKB is for the content that is clearly marked

 Variants curated in OncoKB with an FDA level of evidence

Scope of OncoKB Recognition

- OncoKB's processes for variant evaluation and assertion, data integrity and security, and transparency of evidence
- Policies of oversight and governance
- Processes for ensuring conflicts of interest are minimized and transparent

OncoKB Levels of Evidence Actionable Genes Cancer Genes API / License About News FA BRAF V600E ₪ Oncogenic ⊚ · Gain-of-function ■ · Level 1 ● · Level Dx2 ■ · FDA Level 2 ② Therapeutic Diagnostic FDA-Recognized Content FDA Level 2 Colorectal Cancer FDA Level 2 Anaplastic Thyroid Cancer

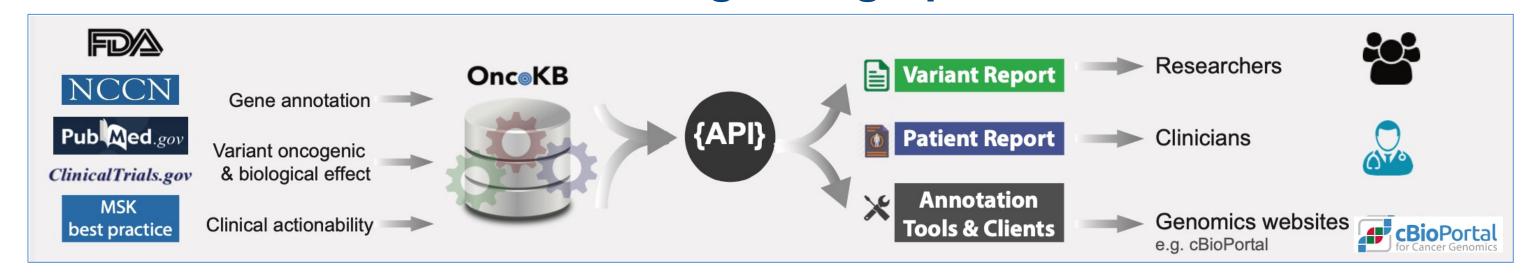
Resistance

11 Genes

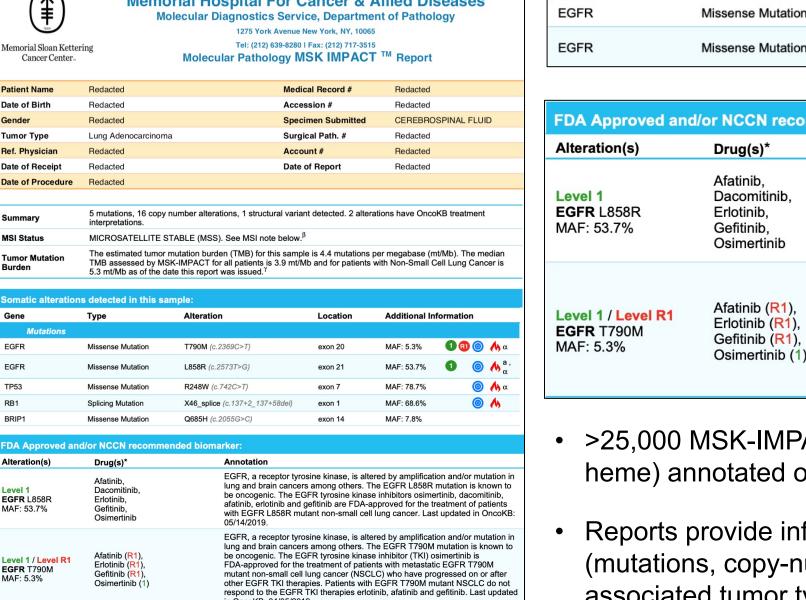
Biological evidence

25 Genes

OncoKB data is available through a high-performance web API



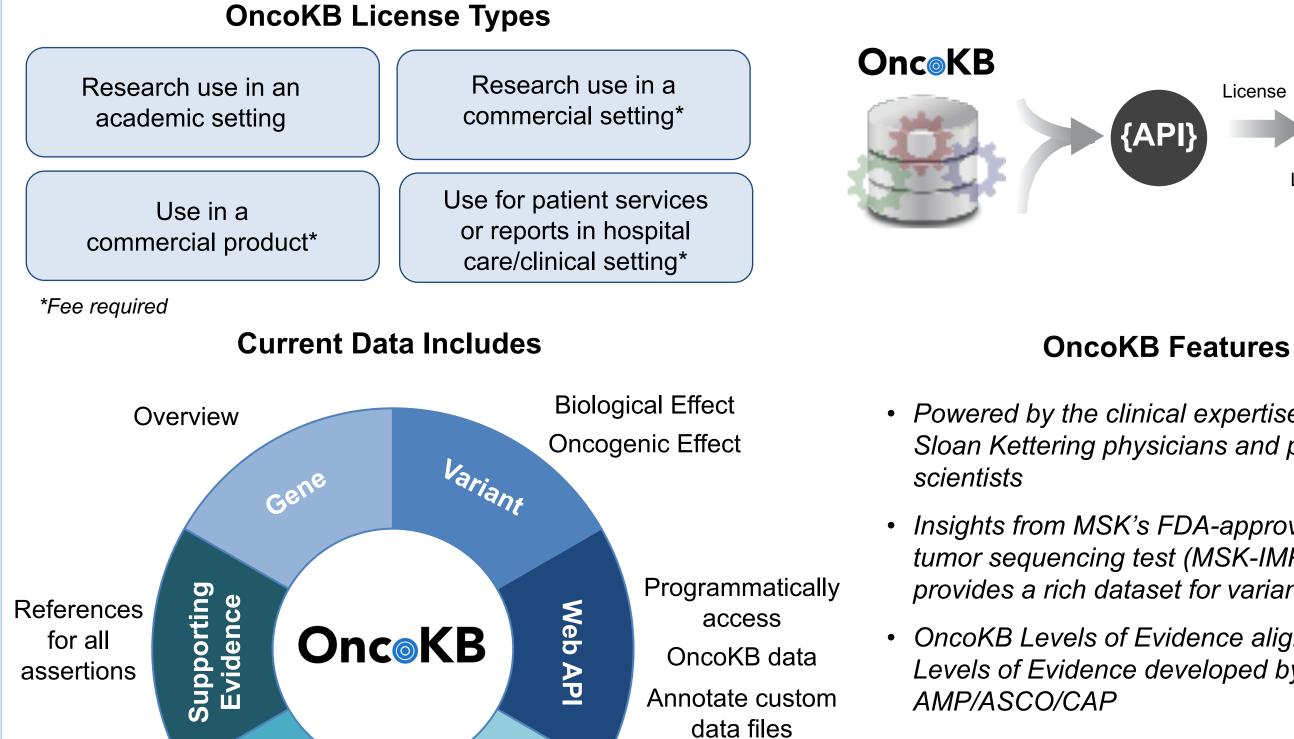
Clinical insight-driven report generation



EGFR	Missense Mutation	L858R (c.2573T>G)	exon 21	MAF: 53.7%		o M
FDA Approved an	nd/or NCCN recommo	ended biomarker:				
Alteration(s)	Drug(s)*	Annotation				
Level 1 EGFR L858R MAF: 53.7%	Afatinib, Dacomitinib, Erlotinib, Gefitinib, Osimertinib	lung and brain o be oncogenic. T afatinib, erlotinib	EGFR, a receptor tyrosine kinase, is altered by amplification and/or mutation in lung and brain cancers among others. The EGFR L858R mutation is known to be oncogenic. The EGFR tyrosine kinase inhibitors osimertinib, dacomitinib, afatinib, erlotinib and gefitinib are FDA-approved for the treatment of patients with EGFR L858R mutant non-small cell lung cancer. Last updated in OncoKB 05/14/2019.			
Level 1 / Level R1 EGFR T790M MAF: 5.3%	Afatinib (R1), Erlotinib (R1), Gefitinib (R1), Osimertinib (1)	lung and brain of the oncogenic. To the concept of the control of	GFR, a receptor tyrosine kinase, is altered by amplification and/or mutation in any and brain cancers among others. The EGFR T790M mutation is known to e oncogenic. The EGFR tyrosine kinase inhibitor (TKI) osimertinib is DA-approved for the treatment of patients with metastatic EGFR T790M nutant non-small cell lung cancer (NSCLC) who have progressed on or after ther EGFR TKI therapies. Patients with EGFR T790M mutant NSCLC do not espond to the EGFR TKI therapies erlotinib, afatinib and gefitinib. Last update of OncoKB: 04/05/2019.			

- >25,000 MSK-IMPACT clinical sequencing reports (solid tumors and heme) annotated over the last two years.
- Reports provide information about the known oncogenicity of all variants (mutations, copy-number alterations and gene fusions), as well as their associated tumor type specific clinical implications.

Licensed users have access to the OncoKB API

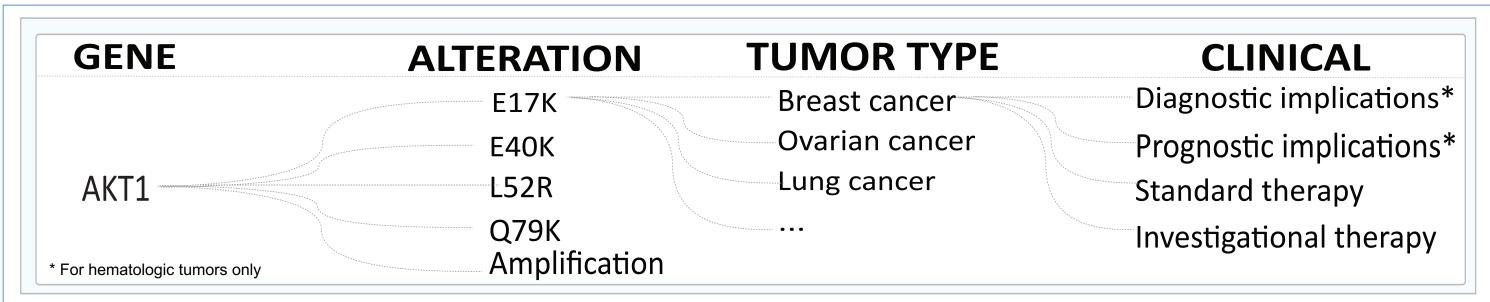


Gene-variant-tumor-type specific drug associations and their corresponding OncoKB level of evidence

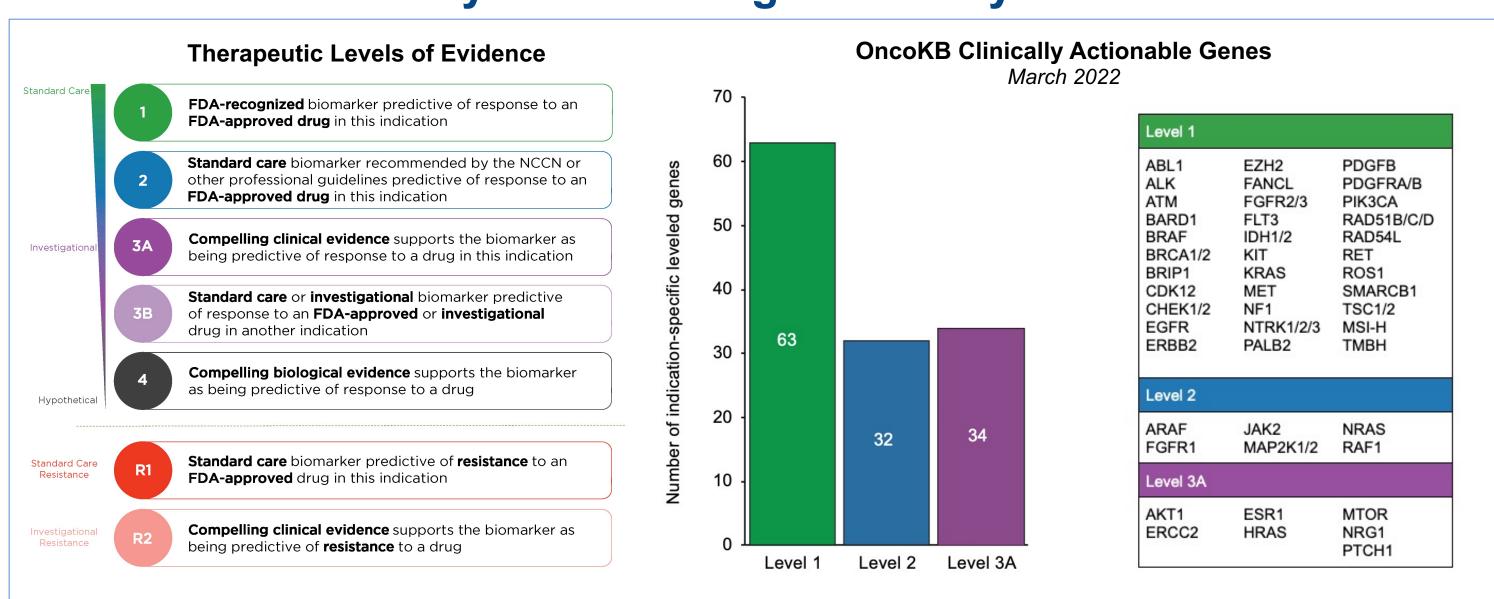


- Powered by the clinical expertise of Memorial Sloan Kettering physicians and physician-
- Insights from MSK's FDA-approved targeted tumor sequencing test (MSK-IMPACT) provides a rich dataset for variant discovery
- OncoKB Levels of Evidence align 1:1 with the Levels of Evidence developed by
- OncoKB annotations can be incorporated into a local instance of cBioPortal for Cancer Genomics providing a custom and dynamic web-based platform for exploring patient sequencing data
- High-performance API for variant annotation
- Support from OncoKB scientists

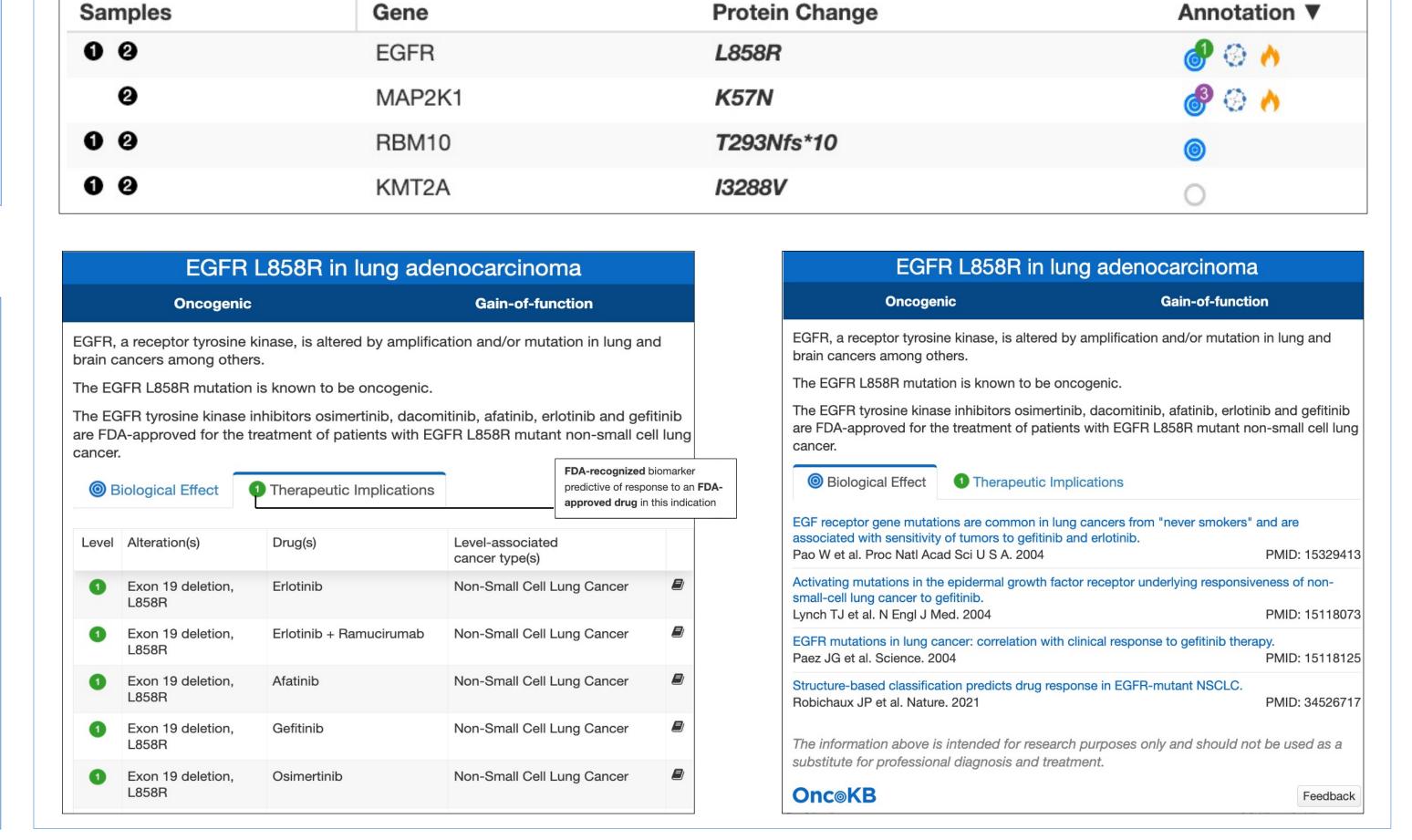
Curation on multiple levels: Gene, Alteration, Tumor Type, MSI/TMB



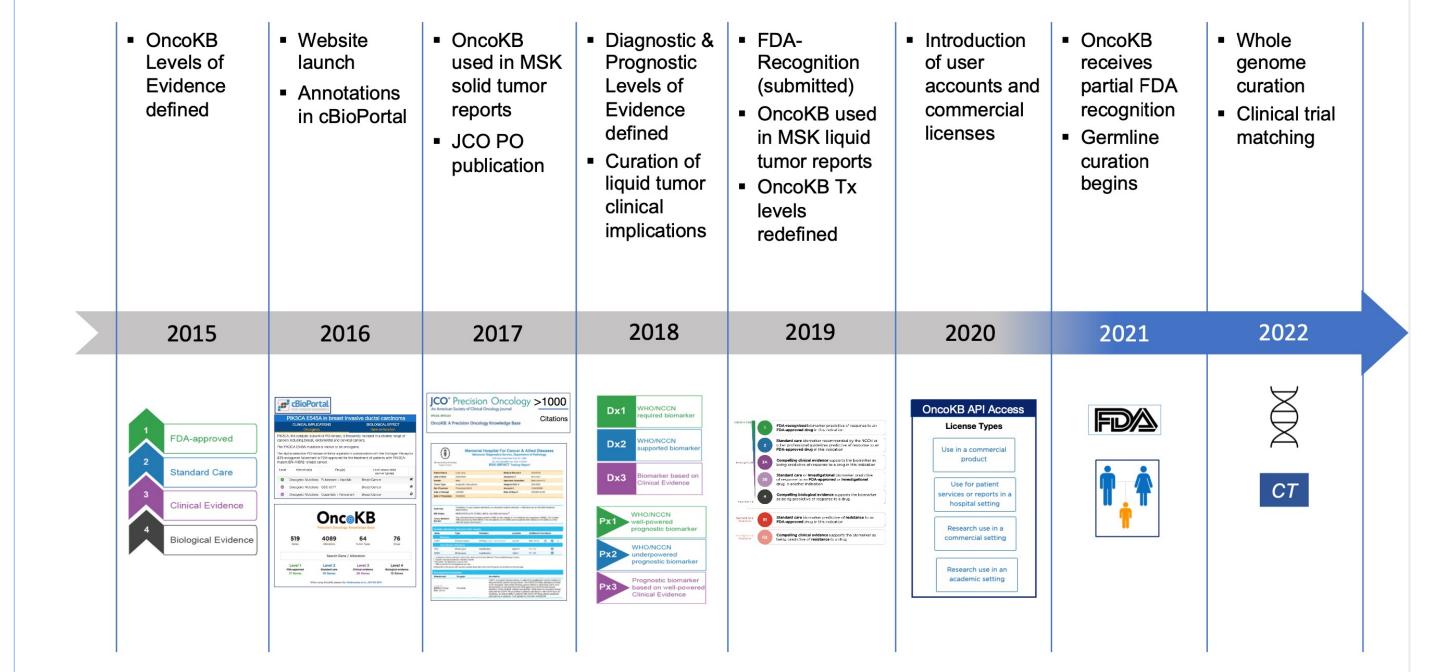
Levels of evidence system for drug sensitivity and resistance



Variant interpretation in cBioPortal



Germline curation, whole genome curation and more coming soon



Resources

Levels 1-4

- OncoKB website: www.oncokb.org
- Licensing information: www.oncokb.org/APIaccess
- Oncotree: <u>www.oncotree.mskcc.org/#/home</u>
- Email us: contact@OncoKB.org

cBioPortal for Cancer Genomics: www.cbioportal.org