Erasmus School of Health Policy & Management

HTA of Cancer Genomics: perspectives transferability and access





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Understanding differences: molecular testing



- Australia (26 Mo people)
- Universal Healthcare (Medicare, tax based) and Private Health Insurance (premium based)
- Public and private hospitals (including pathology labs), specialists work in both
 - Charge service fees to private patients
- Commonwealth manages Medicare: PBS and MBS following rigorous HTA
- State governments fund hospital services through DRGs
 - Complex for interstate patients (e.g. rare cancers like ALL)

- Netherlands (17,6 Mo people)
- Universal Healthcare, Premium based
- Public healthcare (few private centres)
- National Government is responsible for defining benefits scheme (drugs follow HTA process)
- Health insurers (non-for profit) legally bound to offer benefits scheme.
 Competition on premium
- Hospital services are funded through a bundled payment (DOT), including (complex) diagnostics

Implementation of co-deps: item descriptors

Medicare Benefits Schedule - Item 73341

Search Results for Item 73341

Category 6 - PATHOLOGY SERVICES

P7 - Genetics

73341 🛈

Fluorescence in situ hybridisation (FISH) test of tumour tissue from a patient with locally advanced or metastatic non-small cell lung cancer, which is of non-squamous histology or histology not otherwise specified, with documented evidence of anaplastic lymphoma kinase (ALK) mmunoreactivity by immunohistochemical (IHC) examination giving a staining intensity score > 0, and with documented absence of activating mutations of the epidermal growth factor receptor (EGFR) gene, requested by a specialist or consultant physician, to determine:

a. if requirements relating to ALK gene rearrangement status for access to an anaplastic lymphoma kinase inhibitor under the Pharmaceutical Benefits Scheme are fulfilled; or

b. if requirements relating to ALK status for access to pembrolizumab under the Pharmaceutical Benefits Scheme are fulfilled.

Fee: \$400.00 Benefit: 75% = \$300.00 85% = \$340.00

Previous - Item 73340

Next - Item 73342 >



NSCLC treatment landscape & MBS items





Whole-Genome Sequencing to replace SoC tests?



van de Ven et al, J. Molecular Diagnostics, 2021



Cost-effectiveness acceptability of WGS

1.00



Simons et al, 2022



CGS – Actionable Findings (n=217 / 390+ genes)



Equitable access and oucomes



"Mapping disparities in functional outcome of prostate cancer patients" Kendrick Koo et al, BMC Cancer 2022



Mapping of clinical trial participation weighted for cancer incidence. Nancy Tran et al, 2020

Systems models to simulate access to precision oncology



Patient volume ● 0-1500 ● 1501-3000 ● 3001-4500 ● 4501-6000 ● > 6000

Molecular diagnostics



First treatment episode (per hospital)



Implementing Whole-Genome Sequencing







Implementation considerations in HTA

Transferability to the community

- Prevalence (and budget impact) of specific biomarkers (e.g. NTRK, MSI)
 - Depends on testing platform, tested population (e.g. indiginous), cancer stream
- Actionability and incremental benefit of targeted treatment (indirect evidence)
 - Level of evidence of predictive value of co-dep
- Local settings: biopsy (tissue, liquid), sequencing, curation and interpretation
- Implementation issues
 - Sample collection (fresh) and transportation to centres of excellence
 - Data analysis, curation and interpretation vs. treatment in local setting
 - Block vs. fee for service funding? Inter-state patients with different funding mechanisms (state, federal)
- Health systems models to determine effect of system constraints in implementation (granularity of data for policy vs. clinical decisions)