

Hvordan screener man børn for kræft?



Ulrik Kristoffer Stoltze, MD, PhD
om PREDiPOSED projektet

Klinisk genetik

Klinisk pædiatrisk onkogenetik

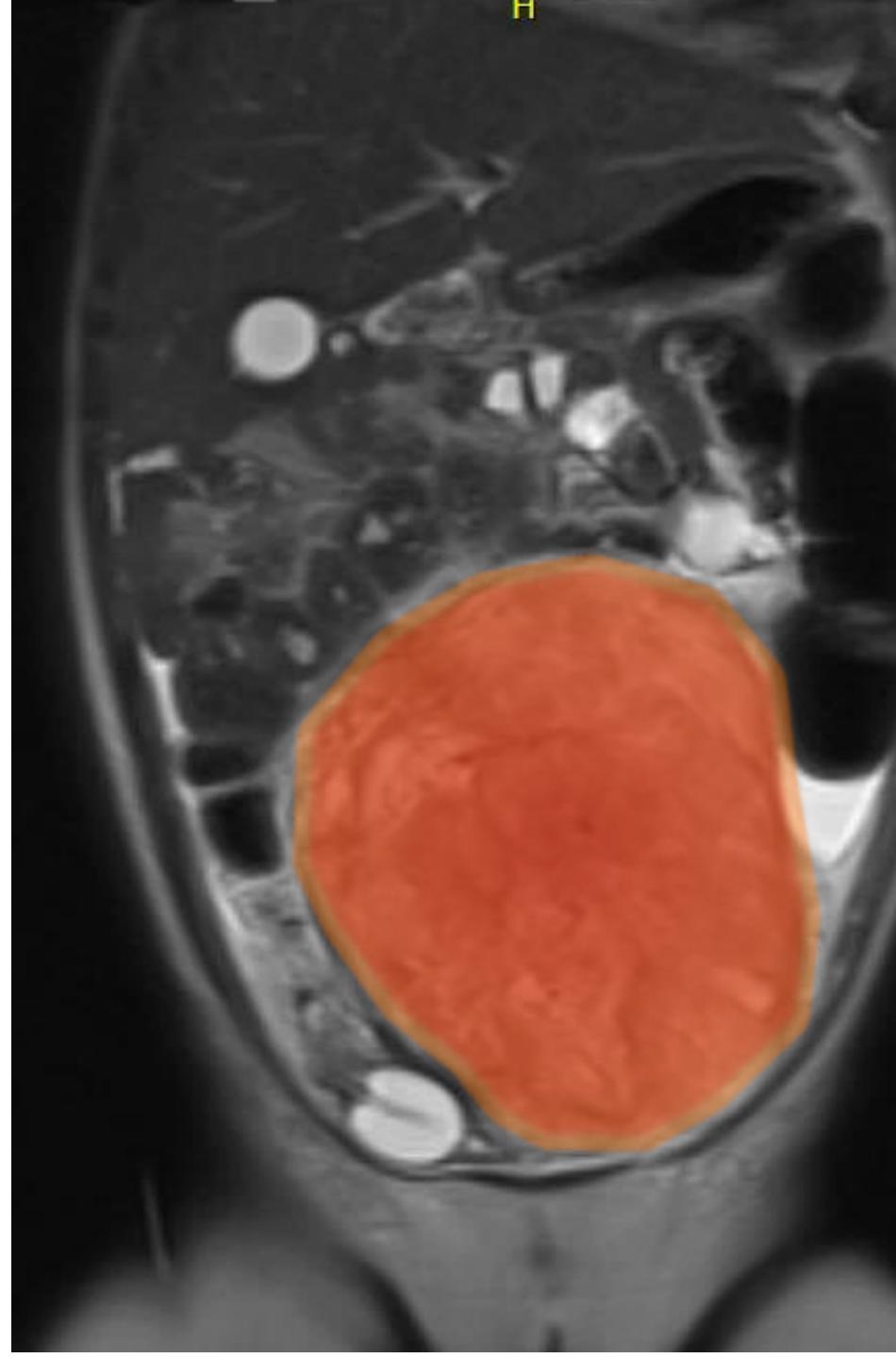




Li-Fraumeni Syndrom

H

Symtomer: måneder







Starter screening

Li-Fraumeni Syndrom

B.T.

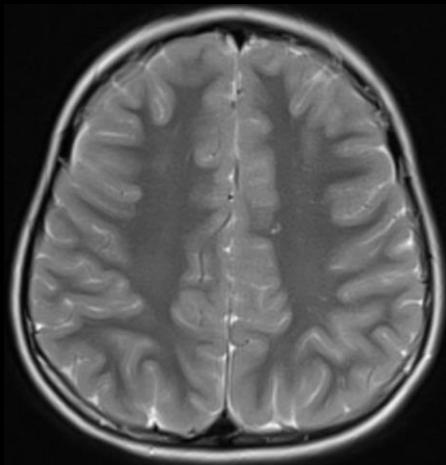
**Ny klinik på Rigshospitalet: Skal
opdage kræft ved børn meget tidligere**



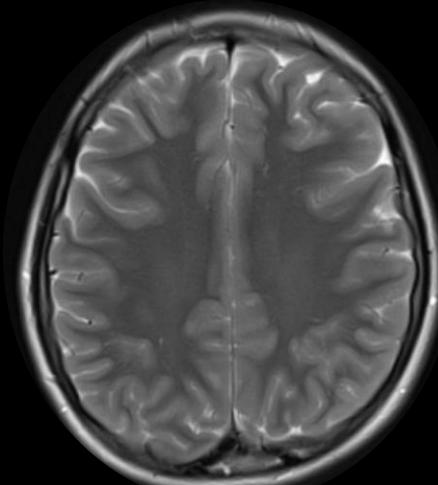


Helkrops MR skanning

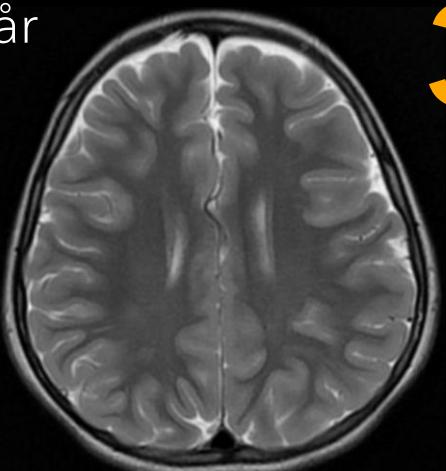
Baseline Helkrops MR skanning på raske børn



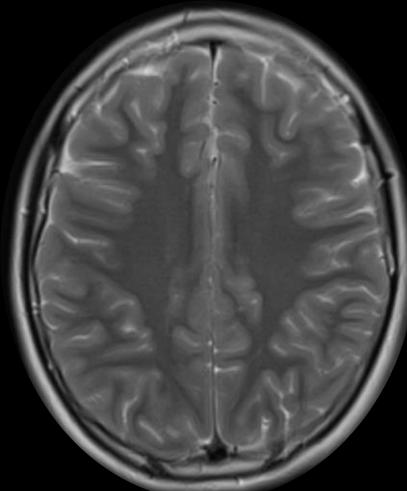
1 Dreng på 9 år
uden fund



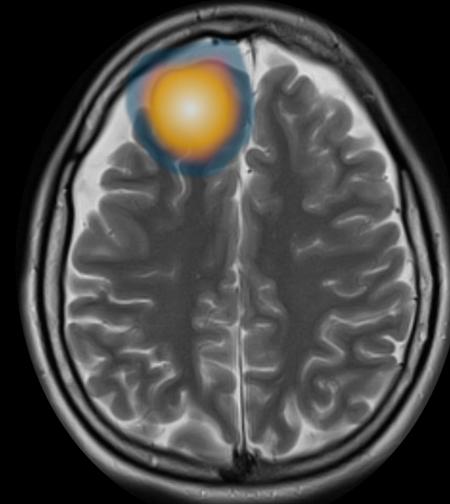
3 Pige på 12 år
uden fund



2 Pige på 8 år
uden fund

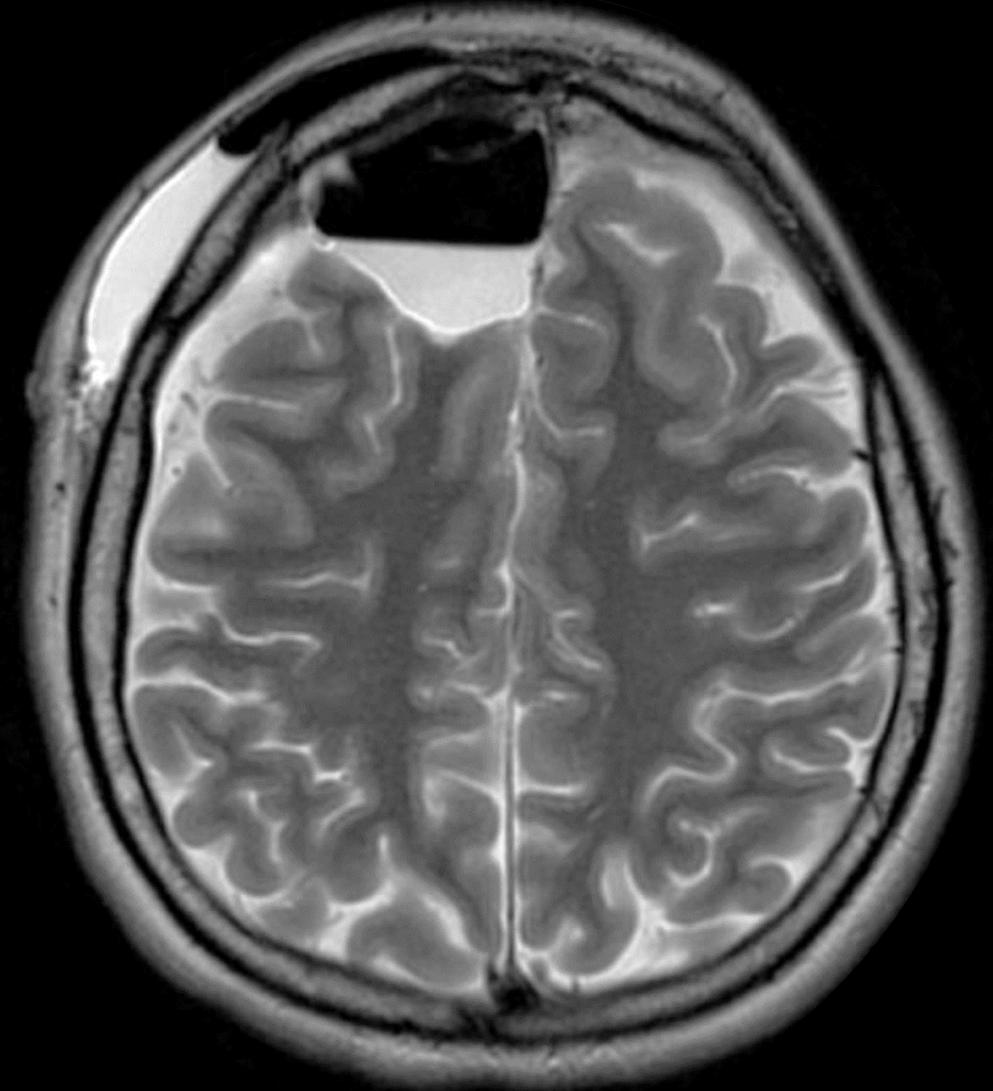


4 Dreng på 13 år
uden fund



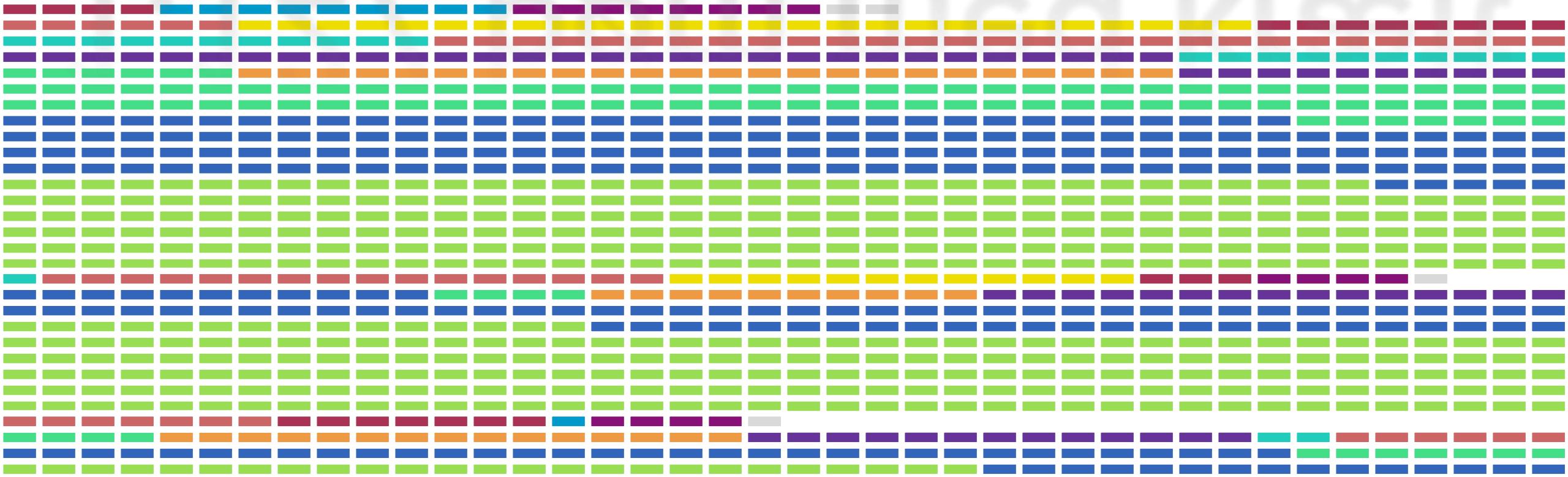
5 Pige på 15 år
tumor

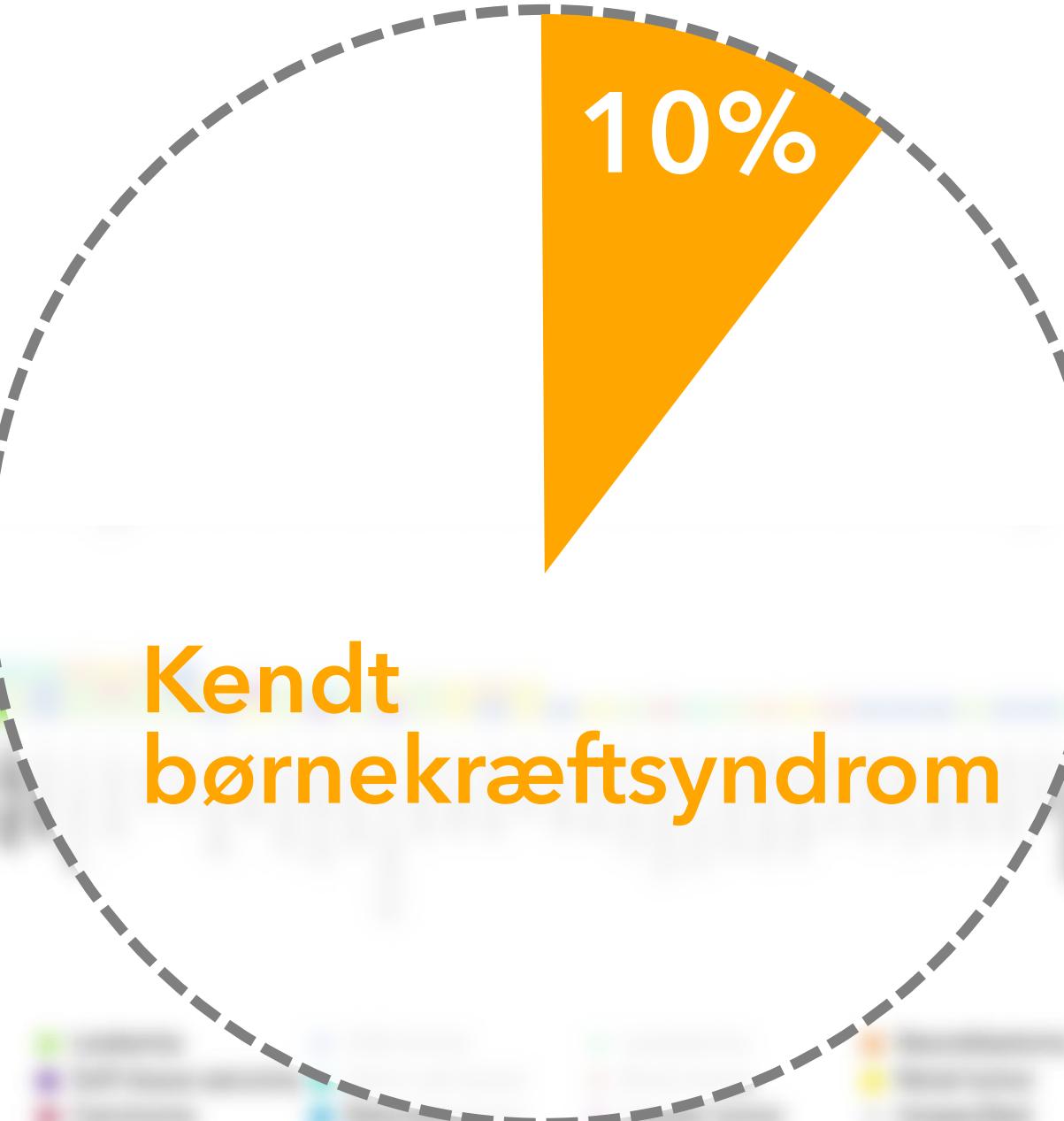
Patient
Sund & rask
Grad 3 astrocytom
Operabel





1.127 børn med kræft



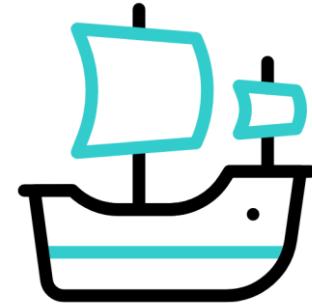


10%

Kendt
børnekræftsyndrom

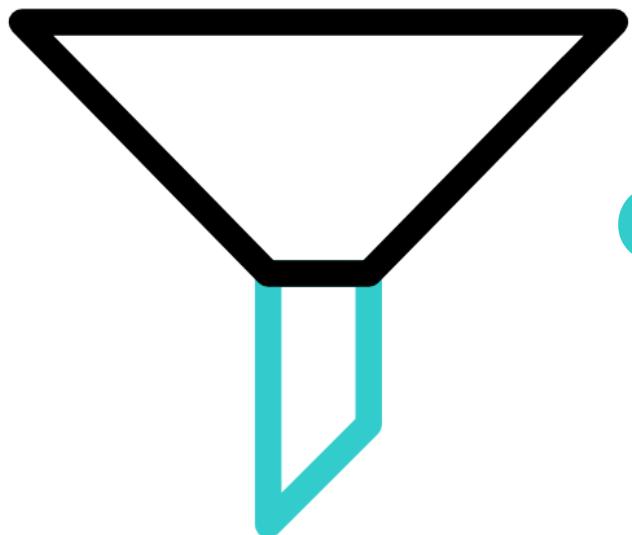


Bombefly &
Børnekæft



Kaptajn Grant
Problemet

4.000.000

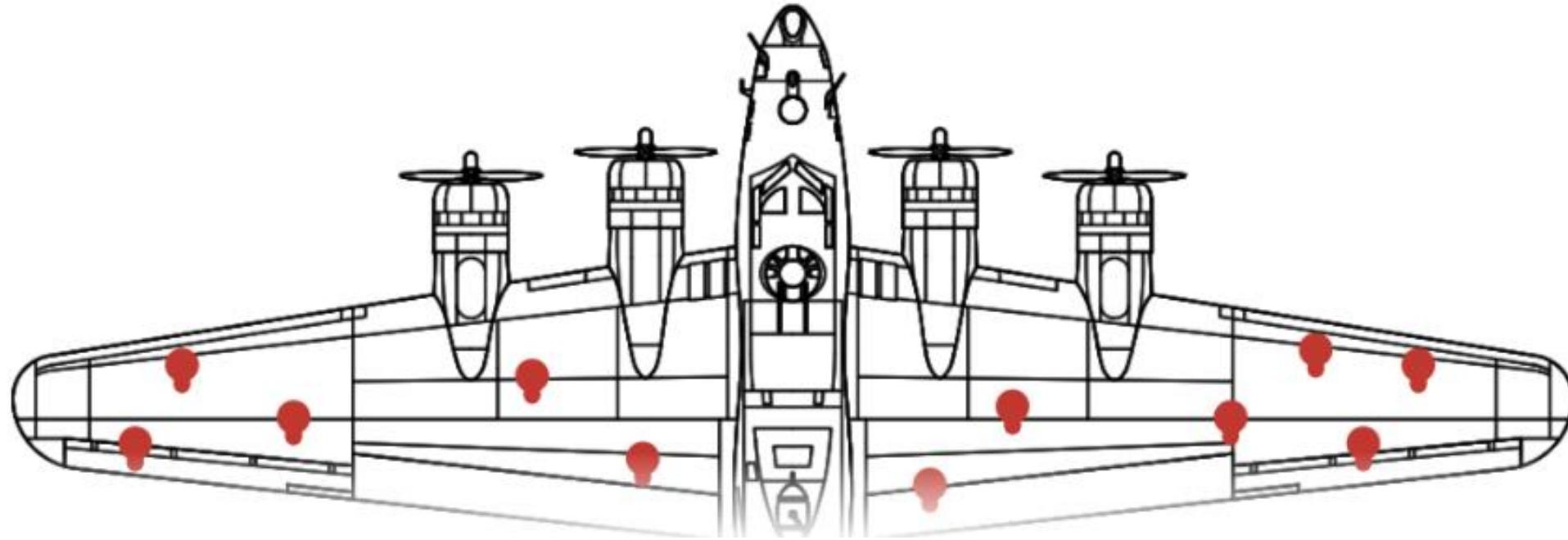


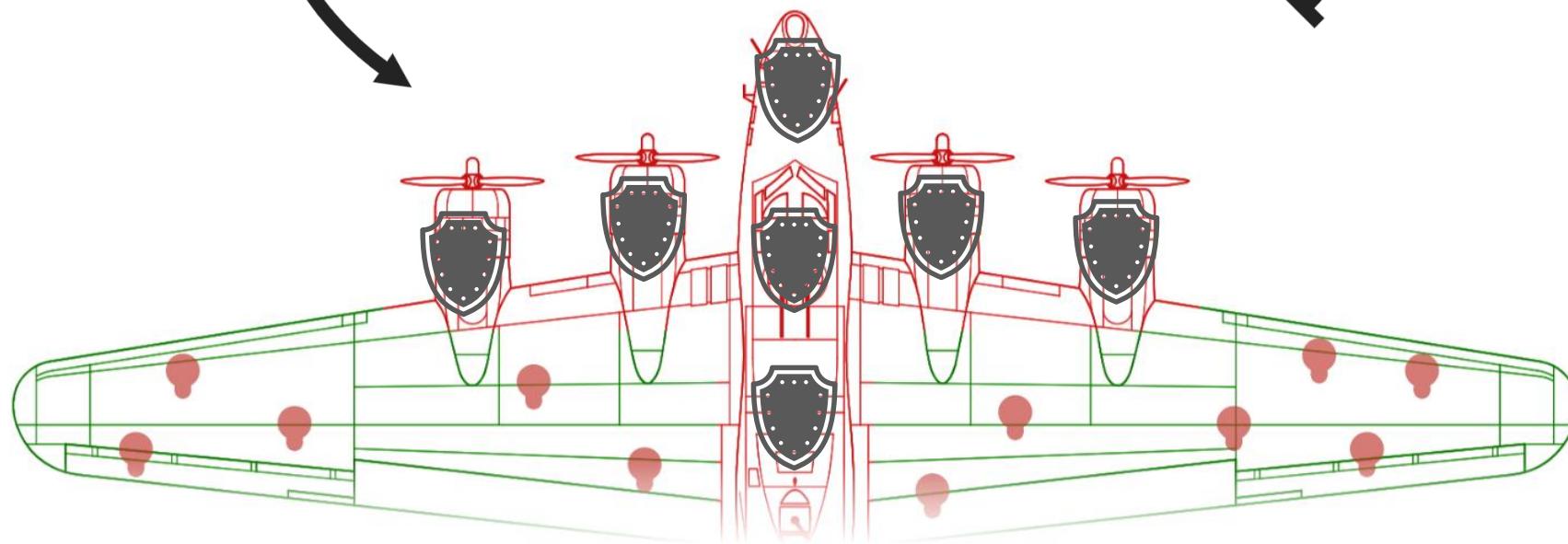
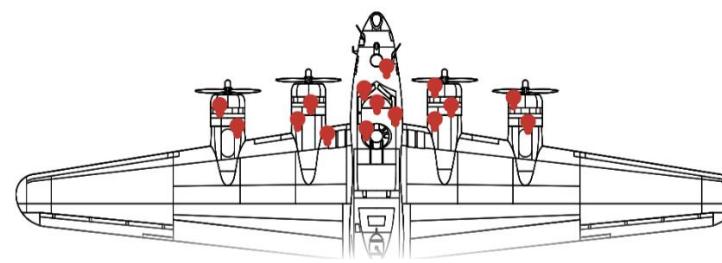
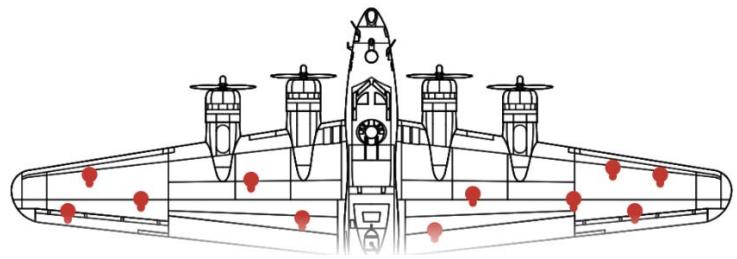
99.99999%

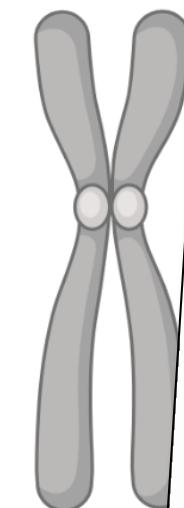
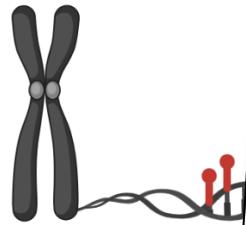
20

A wide-angle photograph of a calm ocean under a bright blue sky with scattered white clouds. The horizon is visible in the distance, and the water is a deep, clear blue.

99.99999%







<https://doi.org/10.1038/s41467-024-45975-9>

The evolutionary impact of childhood cancer on the human gene pool

Received: 16 March 2023

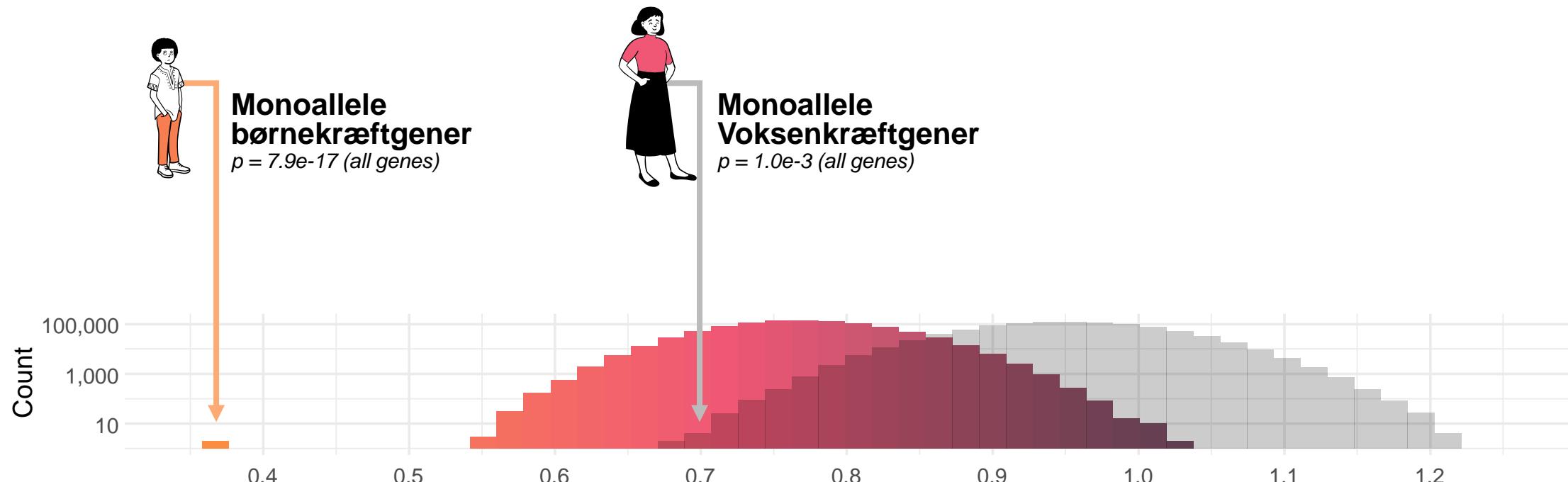
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Check for updates

Ulrik Kristoffer Stoltze ^{1,2,3}, Jon Foss-Skiftesvik ^{1,4}, Thomas van Overeem Hansen ^{2,5}, Simon Rasmussen ^{6,7}, Konrad J. Karczewski ^{3,7,8,9}, Karin A. W. Wadt ^{2,5} & Kjeld Schmiegelow ^{1,5}

Germline pathogenic variants associated with increased childhood mortality must be subject to natural selection. Here, we analyze publicly available germline genetic metadata from 4,574 children with cancer [11 studies; 1,083 whole exome sequences (WES), 1,950 whole genome sequences (WGS), and 1,541 gene panel] and 141,456 adults [125,748 WES and 15,708 WGS]. We find that pediatric cancer predisposition syndrome (pCPS) genes [$n = 85$] are highly constrained, harboring only a quarter of the loss-of-function variants that would be expected. This strong indication of selective pressure on pCPS genes is found across multiple lines of germline genomics data from both pediatric and adult cohorts. For six genes [*ELP1*, *GPR161*, *VHL* and *SDHA/B/C*], a clear lack of mutational constraint calls the pediatric penetrance and/or severity of associated cancers into question. Conversely, out of 23 known pCPS genes associated with biallelic risk, two [9%, *DIS3L2* and *MSH2*] show significant constraint, indicating that they may monoallelically increase childhood cancer risk. In summary, we show that population genetic data provide empirical evidence that heritable childhood cancer leads to natural selection powerful enough to have significantly impacted the present-day gene pool.



Højt
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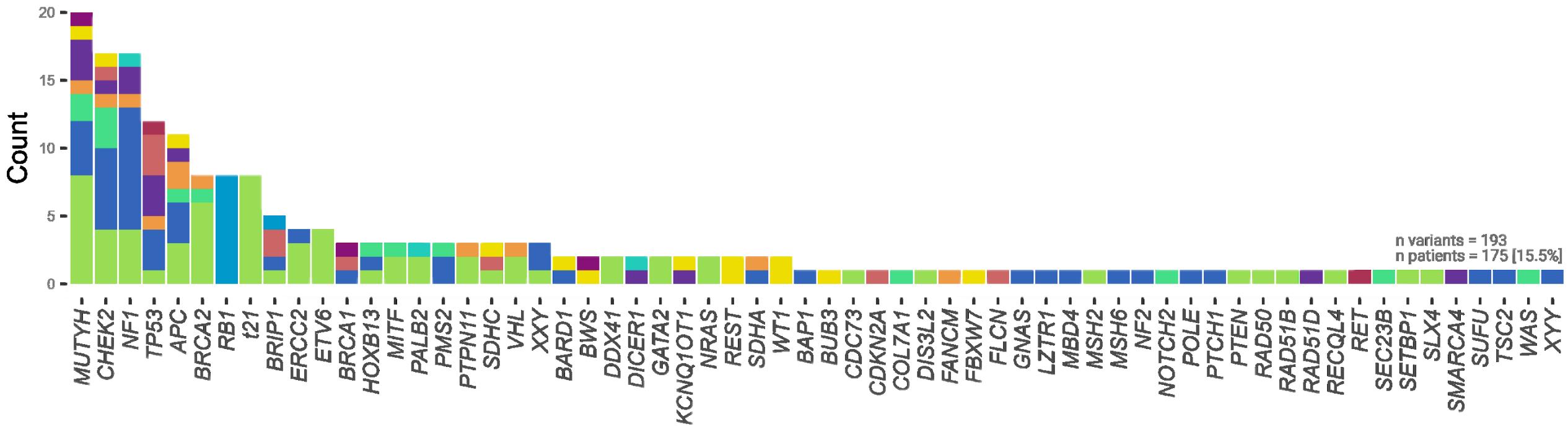
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Thomas van Overem Hansen ^{2,5}, Simon Rasmussen ^{3,6,7},
Konrad J. Karczewski ^{3,7,8,9}, Karin A. W. Wadl ^{2,3} & Kjeld Schmiegelow ^{1,5}

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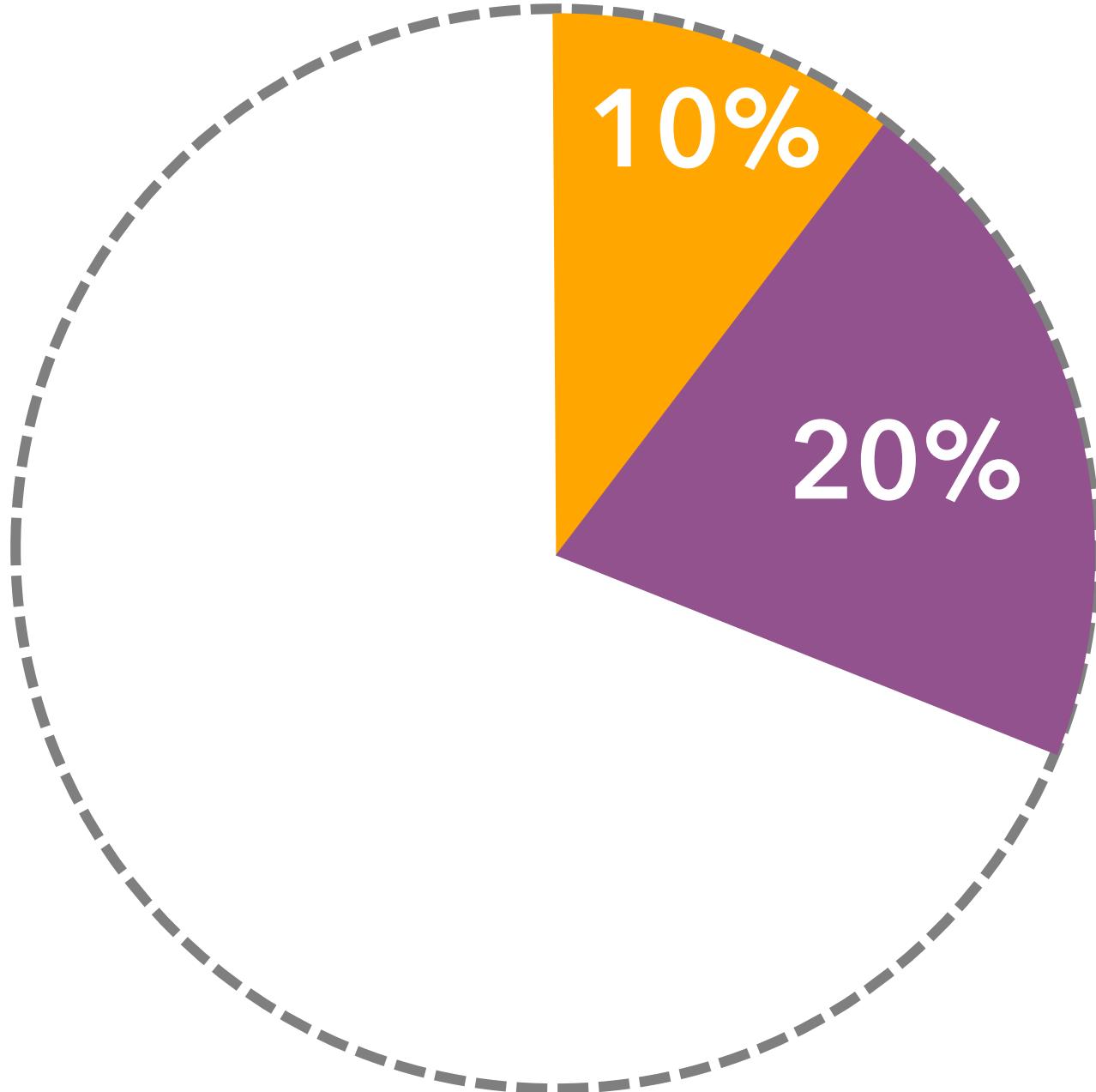
A wide-angle photograph of a calm ocean under a bright blue sky with scattered white clouds. The horizon is visible in the distance, and the water is a deep, clear blue.

99.99999%

Børn med kræft har
en overhyppighed af
evolutionært atypiske
genforandringer



Børn med k~~raft~~^{raft} har
en overhyppighed af
evolutionært atypiske
Ugørkforandringer

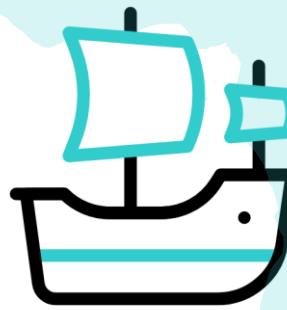


Kendt
børnekræftsyndrom

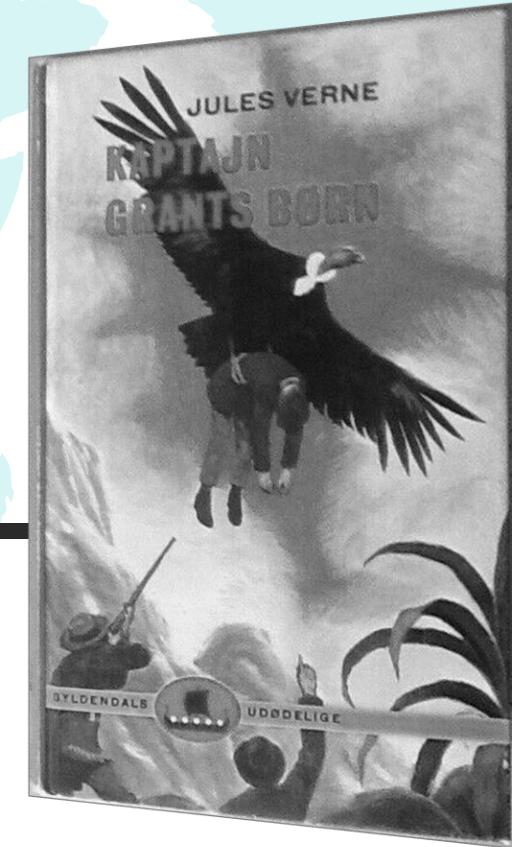
Endnu ukendt
børnekræftsyndrom?

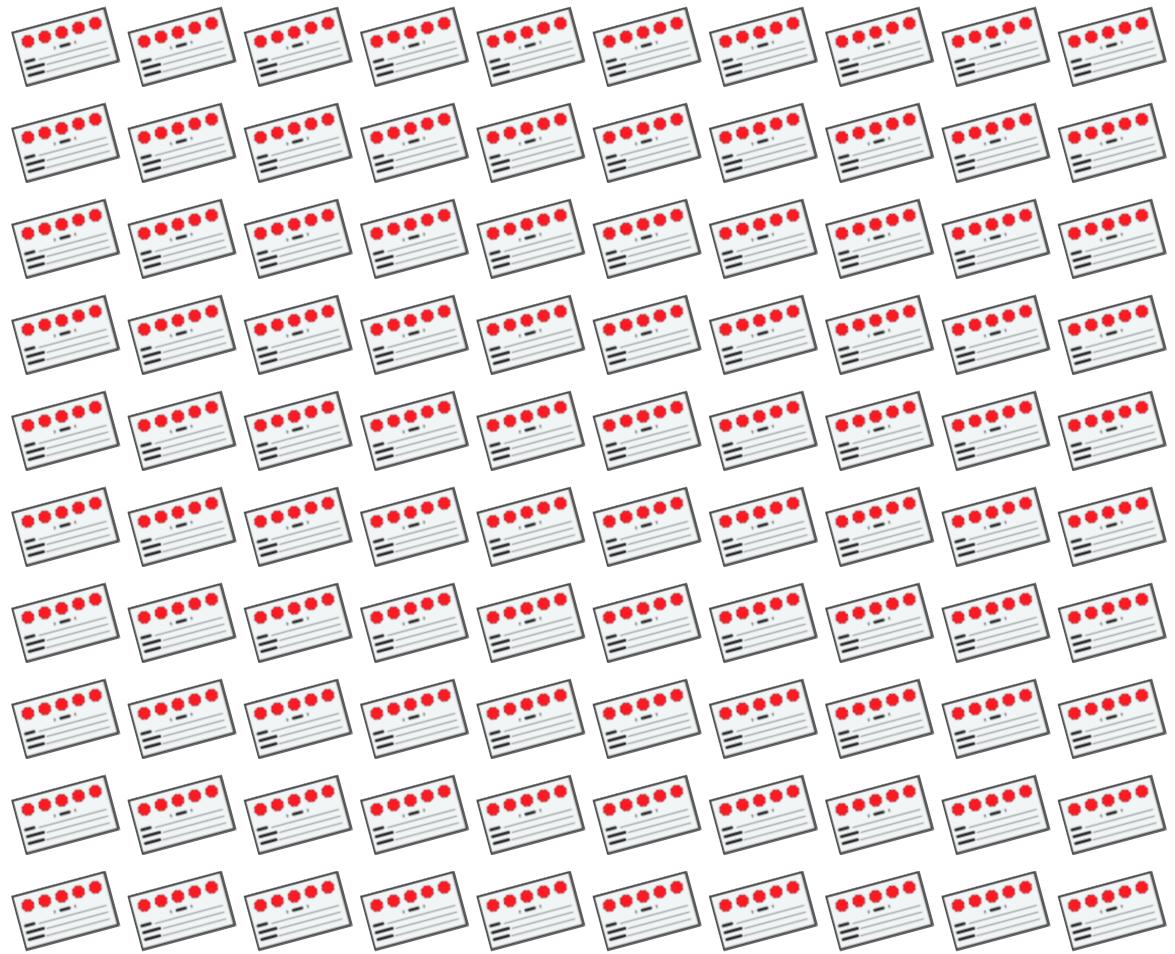


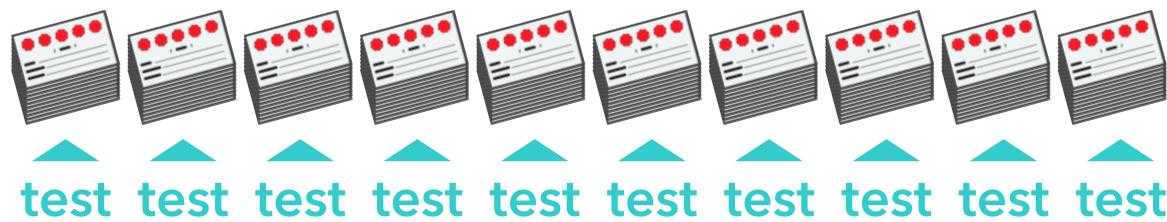
Bombefly &
Børnekæft



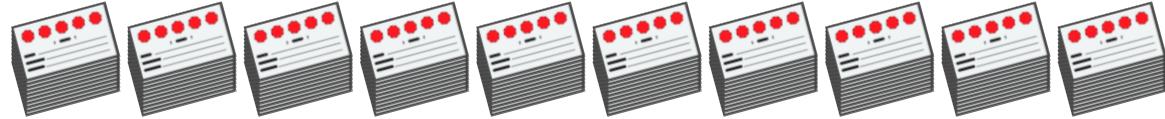
Kaptajn Grant Problemet

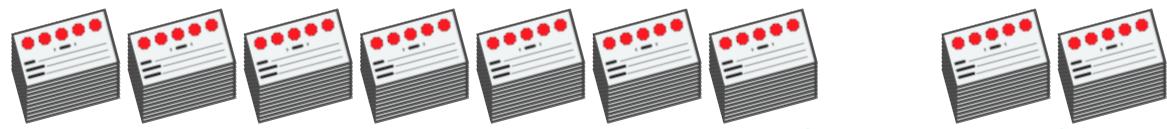






TP53 mutation





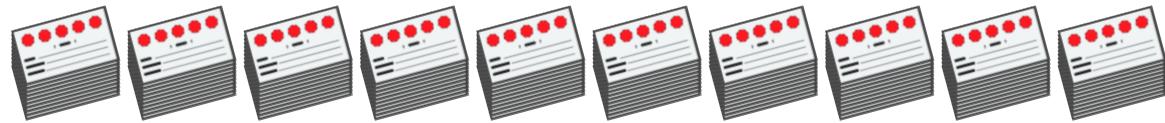
TP53 mutation?



TP53 mutation

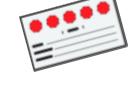
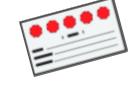
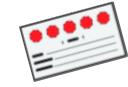
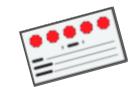
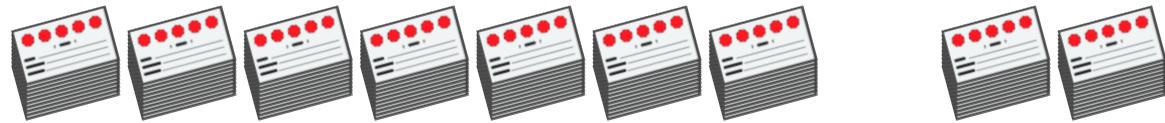


TP53 mutation



TP53 mutation ➤

TP53 mutation

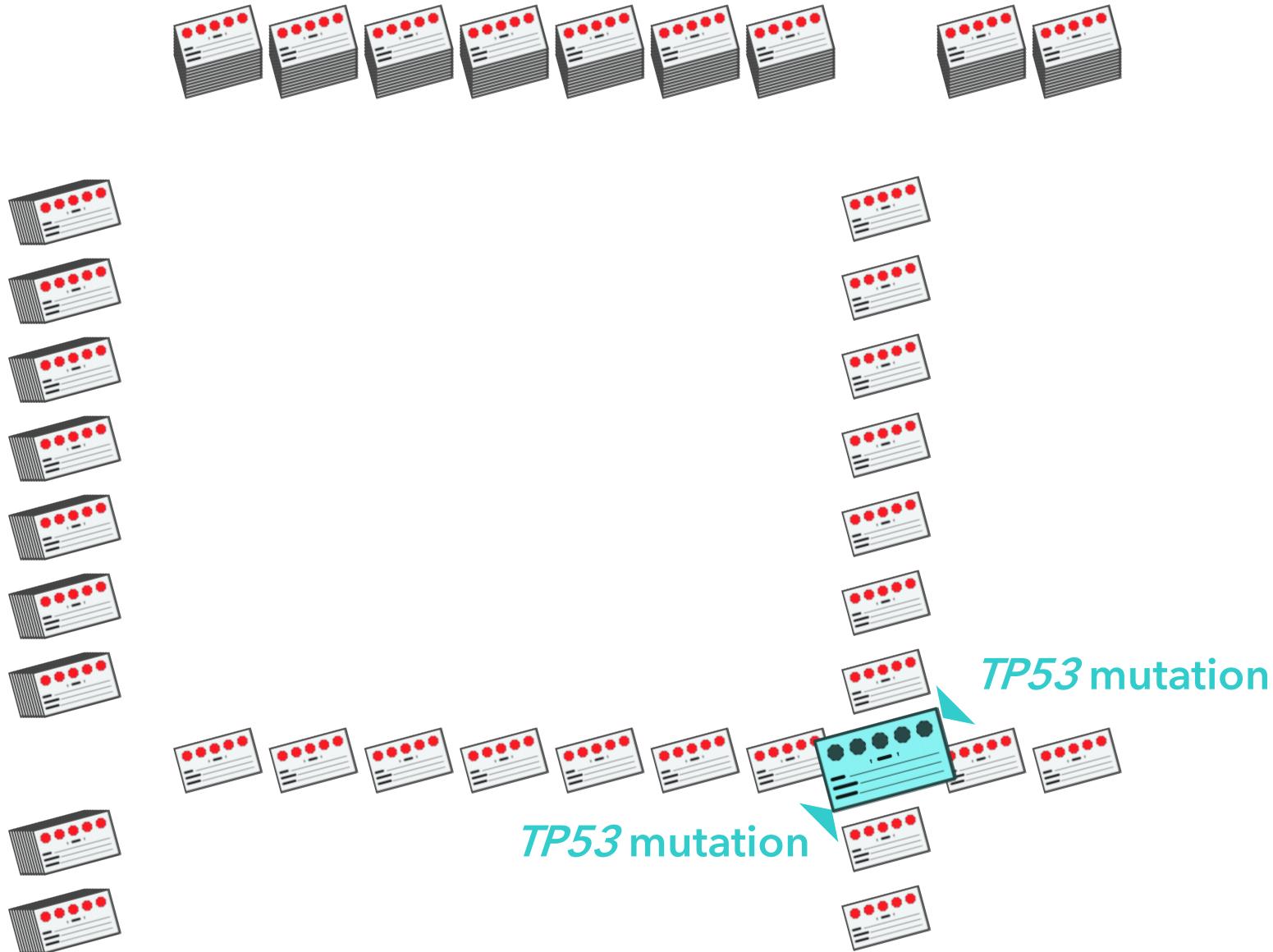


TP53 mutation



TP53 mutation





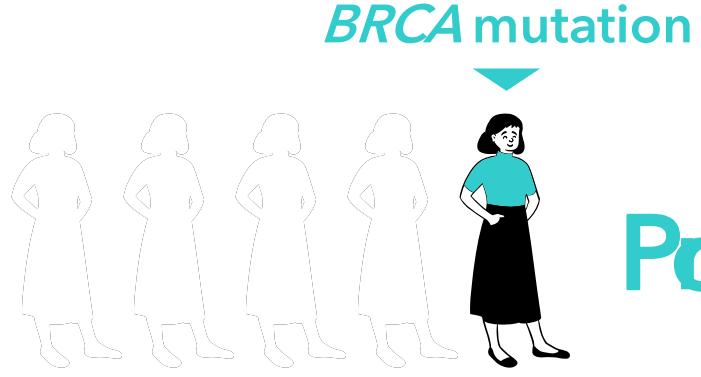


Diagnose



Omkostninger

60.000 nyfødte / år
~15 er som Vega
↓>50% børnekraeft-dødelighed



Pøførlitøasscreening



Sjældne sygdomme



PREDiSPOSED

Population-based Retro- & prospective Evaluation
of Diagnostic Sequencing for Pediatric & Oncogenetic Syndromes' Early Detection

STOR TAK TIL...



Kjeld
Schmieelow



Karin
Wadt



Thomas
v. Overeem Hansen



Simon
Rasmussen



Henrik
Hjalgrim



Jonas
Bybjerg-Grauholt



Christian
Munch Hagen



Allan
Lund



Signe Holst
Søegaard



Ayo
Wahlberg



Jacob Tfelt-
Hansen



Tania
Masmas



Lene
Rottensen

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Vera Rasmussen
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Bettina Lundgren
Mads Melbye
Charlotte K. Lautrup
Astrid Saksager

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Daniel Dybdal
Ayo Whalberg
Anne-Marie Gerdes
Jane Frederiksen
Rene Mathiasen
Anja Hirche
Solvej Kullegaard
Astrid Saksager

Friends and family!

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Christian Hagen
Henrik Hjalgrim
Marie Bækvd-Hansen

International advisory board:

Michael Walsh
Christian Kratz
Nick Hayward

Bonkolab PhDs, postdocs, nurses, and all staff

THE PATIENTS AND THEIR FAMILIES



FUNDING

børne | cancer | fonden



Innovation Fund Denmark

Interreg
EUROPEAN UNION

