

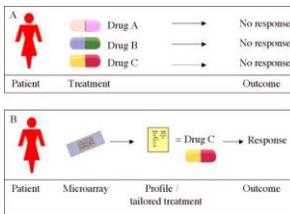


Molekulare Pathologie in der modernen Diagnostik des Gastrointestinaltraktes

Prof. Dr. med. Luigi Tornillo
GILAB AG
XXII. Diagnostik Symposium
Schaan, 10.03.2016

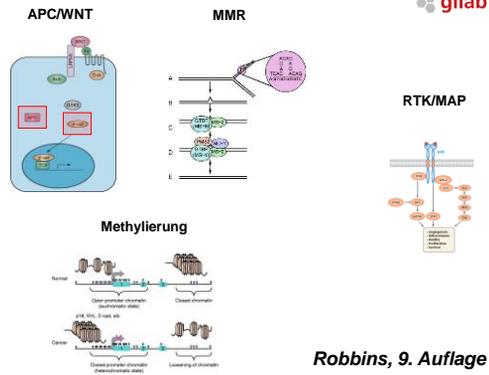


- Einleitung
- Technik
- Anwendungen
- NGS
- Allgemeine Schlussfolgerungen

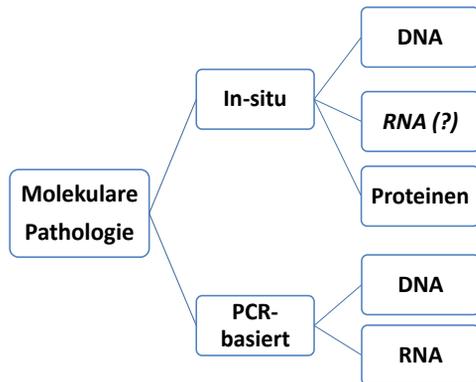


“The right drug, the right dose, for the right patient, at the right time”

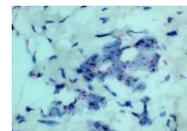
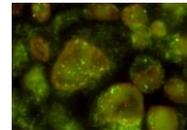
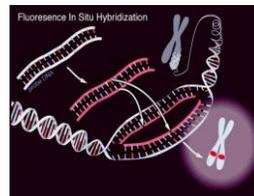
Brown et al. 2003

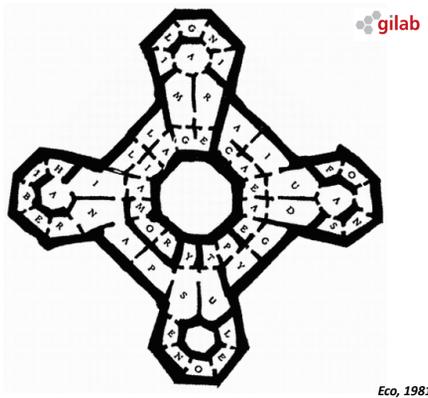
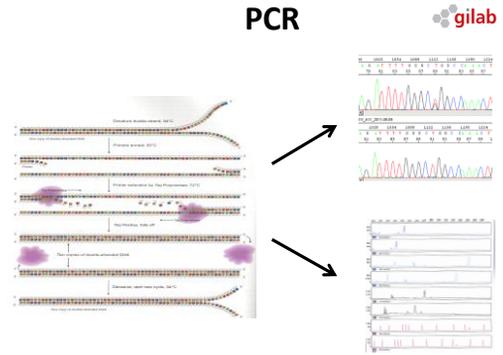
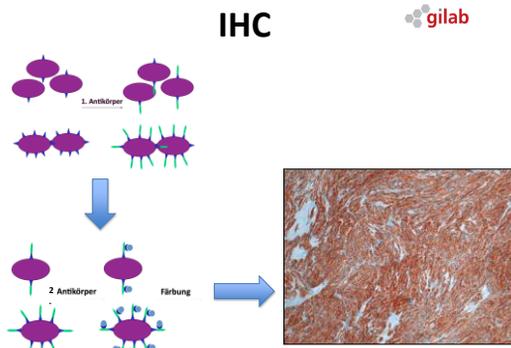


Robbins, 9. Auflage



In-situ



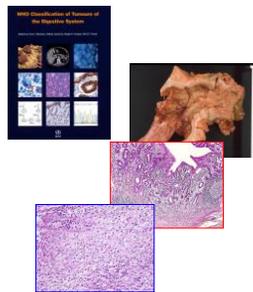


Magenadenokarzinom

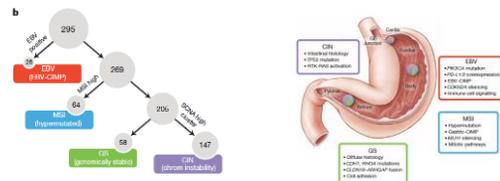
WHO 2010

Tab. 2 Klassifikation der Weltgesundheitsorganisation der Karzinome des Magens [13]

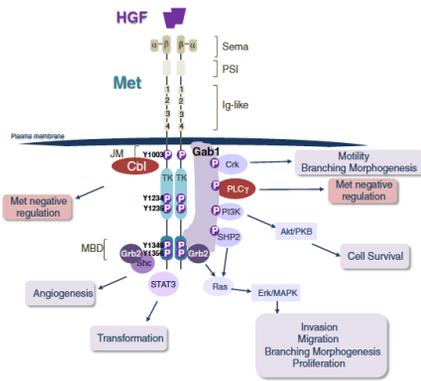
Epitheliale Tumoren	ICD-O M
Adenokarzinom	8140/3
1. Papilläres Adenokarzinom	8260/3
2. Tubuläres Adenokarzinom	8211/3
3. Muzinöses Adenokarzinom	8480/3
4. Gering kohäsives Karzinom (einschließlich Siegelringzellkarzinom und anderer Varianten)	8490/3
5. Gemischtes Adenokarzinom	8255/3
Adenosquamoses Karzinom	8560/3
Karzinom mit lymphoidem Stroma (medulläres Karzinom)	8512/3
Hepatoides Adenokarzinom	8576/3
Plattenepithelkarzinom	8070/3
Undifferenziertes Karzinom	8020/3



Magenadenokarzinom: Molekulare Klassifikation



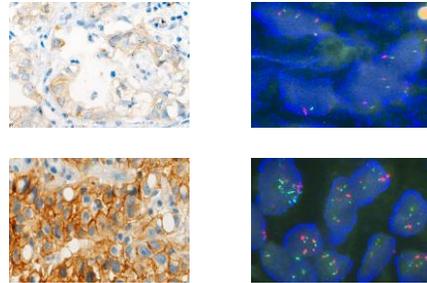
The Cancer Genome Atlas Research Network, Nature, 2014



CMET

IHC

FISH



„Eine (neo)adjuvante Therapie mit zielgerichteten Substanzen alleine oder in Kombination mit Chemotherapie soll außerhalb von Studien **nicht** durchgeführt werden“

S-3 Leitlinien zum Magenkarzinom, 2013

Schlussfolgerungen I



- ERBB2 (HER2) wichtig für die Progression vom metastat. MK und mit Antwort zur gezielten Therapie assoziiert
- Bestimmung des HER2-Status **wichtig im metastatischen Magenkarzinom**
- Andere RTK in Rahmen von Studien
- Molekulare Klassifikation (?)

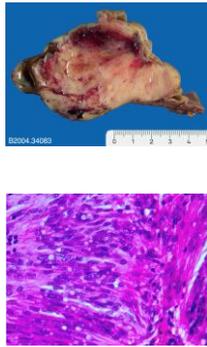


GIST



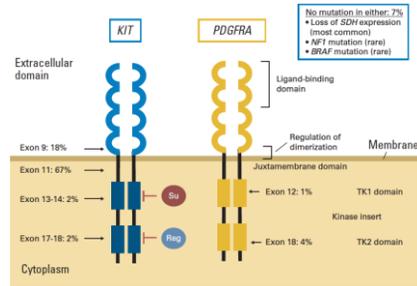
2001

„Primäre spindelzellige epitheloidzellige, gemischte oder pleomorphe mesenchymale Neoplasien des GI-Trakts“



Agaimy & Schneider-Stock, Pathologie 2010

GIST: Pathogenese



Cioffi & Maki, JCO 2015

GIST: Prognose



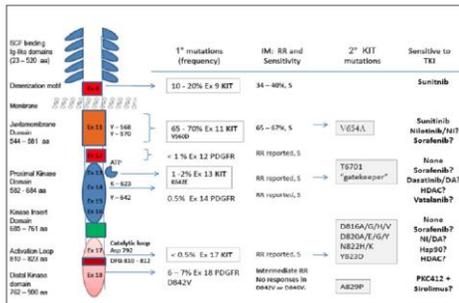
Mitotic Index (per 50 HPF)	Tumor Parameters	Size (cm)	Risk of Progressive Disease* (%)		
			Gastric	Duodenum	Jejunum/Ileum
≤5	≤2	None (0%)	None (0%)	None (0%)	None (0%)
	>2, ≤5	Very low (1.9%)	Low (4.3%)	Low (8.3%)	Low (8.5%)
	>5, ≤10	Low (3.6%)	Moderate (24%)	Moderate (24%)	Moderate (24%)
>5	>10	Moderate (10%)	High (52%)	High (34%)	High (57%)
	≤2	None†	High (73%)	High (50%)	High (54%)
	>2, ≤5	Moderate (16%)	High (85%)	High (50%)	High (52%)
	>5, ≤10	High (55%)	High (80%)	High (80%)	High (71%)
	>10	High (86%)	High (90%)	High (86%)	High (71%)

AFIP, JNCNN, 2007 (sog. "Miettinen's criteria")

GIST: RTK-Mutationen

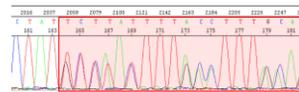
- **Lokalisation**
 - KIT Exon 9 Dünndarm, PDGFRA Magen
 - **Histologie**
 - PDGFRA epitheloid, WT epitheloid
 - **Prognose**
 - KIT exon 9 schlecht (?)
 - KIT exon 11 Deletion schlecht, PDGFRA oft indolent
 - **Therapie - Resistenz**
- **Primäre Resistenz (10-20%)**
 - Ex 9, Ex 13-17, "Wild-Typ"
 - **Sekundäre Resistenz (24 Mo)**
 - Zweite Mutation (Ex 13-17)
 - KRAS?, BRAF?

GIST: RTK-Mutationen

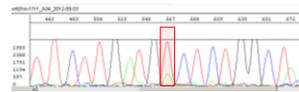


Gounder, Cancer Chemoter Pharmacol 2011

GIST: RTK-Mutationen



p.Ala502-Tyr503dup



p.Asp816Val

Schlussfolgerungen II



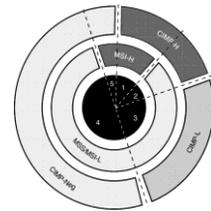
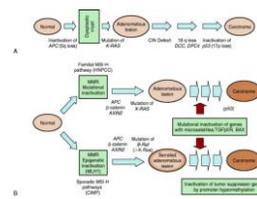
- „Gold-standard“ in GIST ist Chirurgie
- RTKI für adjuvante Therapie
- **Mutationsanalyse immer, wenn adjuvante Therapie angezeigt**
- Die Zukunft: neoadjuvante Therapie ?



KRK



Kolonkarzinom



Robbins, 9. Auflage

Jass, Histopathology, 2006

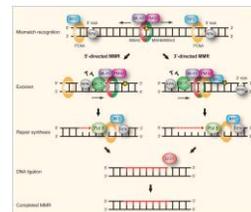
KRK: Molekulare Einteilung

- 3 „grobe“ molekulare „pathways“
 - APC-CIN (Adenom-Karzinom Sequenz, FAP)
 - MSI-CS (Lynch, Epigenetik)
 - CIMP (Epigenetik)
- Prädiktive Faktoren
 - G-proteins, PI3KCA

MMR



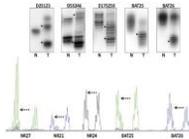
- Ungepaarte Nukleotiden
- Insertion-Deletion loops
- Komplexer „Apparat“



MSI

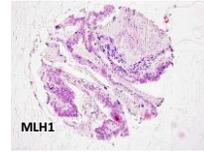
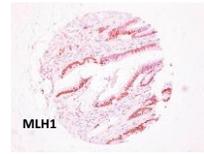


- Kurze Sequenzen ncDNA
 - Mono-, Di-, Trinukleotiden
 - AAAAA..., CACACA..., CAGCAGCAG
- Bethesda Panel
 - 5 Marker
 - ≥ 2 Instabil



MMR

- IHC
 - MLH1, MSH2, MSH6, PMS2
 - Sensitivität 92.5%
 - Spezifität 100%
 - PNW 96.7%
 - PPW 100%



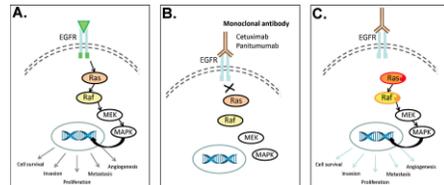
Lindor et al., JCO 2002

MMR IHC: Bedeutung



MLH1	MSH2	MSH6	PMS2		
+	+	+	+	MSS	=
-	+	+	-	MSI	Sporadisch od. Germline
+	-	-	+	MSI	Germline
+	+	-	+	MSI	Germline
+	+	+	-	MSI	Germline

KRK: gezielte Therapie

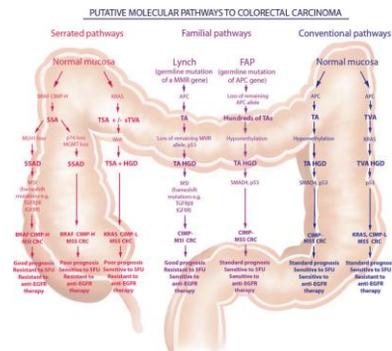
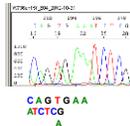
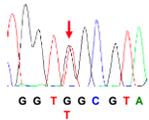


Bedeir, APLM 2013

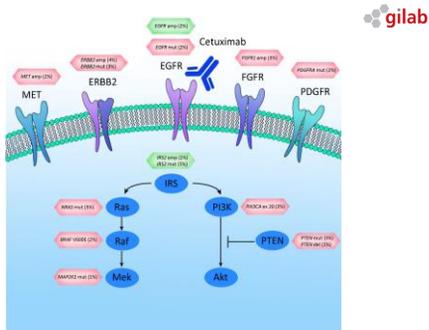
KRK: gezielte Therapie



- RAS
 - 30-40% der Fälle
 - Negativer prediktiver Wert für anti-EGFR
 - Subset prognostischer Wert (G13D) und immunogen
- BRAF
 - 10-15%
 - Negativer prediktiver Wert für anti-EGFR
 - Serrated pathway (MSI sporadisch)
 - Schlechtere Prognose



Bettington, Histopathology, 2013

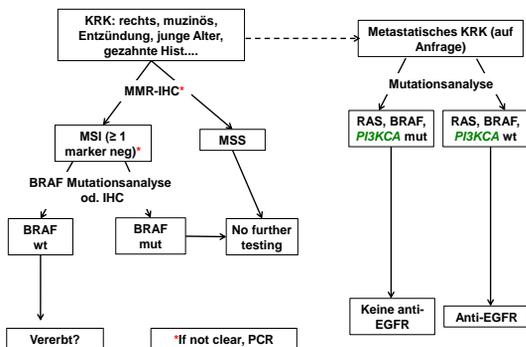


Bertotti et al., Nature 2015

KRK: Molekulare Klassifikation

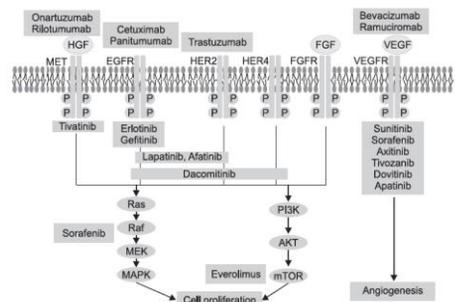
CMS1 MSI immune	CMS2 Canonical	CMS3 Metabolic	CMS4 Mesenchymal
14%	37%	13%	23%
MSI, CIMP high, hypermutation	SCNA high	Mixed MSI status, SCNA low, CIMP low	SCNA high
BRAF mutations		KRAS mutations	
Immune infiltration and activation	WNT and MYC activation	Metabolic deregulation	Stromal infiltration, TGF-β activation, angiogenesis
Worse survival after relapse			Worse relapse-free and overall survival

Guinney et al., Nat Med, 2015



Schlussfolgerungen III

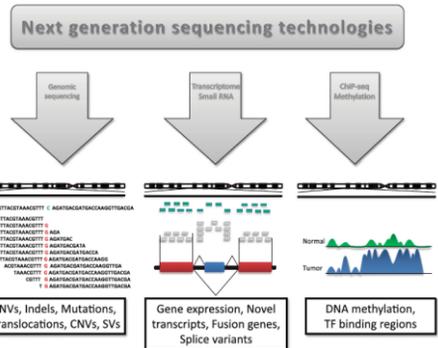
- MSI-H BRAFmut sporadisch
- MSI-H Immunogen
- RAS Mutationen Exonen 2,3,4 prädiktiv (anti-EGFR)
- RAS Mutationen Exone 2,3,4 UND BRAF Mutationen (V600E) prognostisch
- PIK3CA (Exon 9 und 20) prädiktiv (Aspirin UND Anti-EGFR)



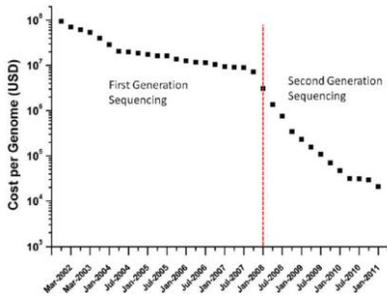


„Normal human genomes are all alike, but every cancer genome is abnormal in its own way.“

Meyerson et al., Nat Rev Gen 2010



Schweiger, Cancer Met Rev 2010



Niedringhaus et al., Anal Chem 2009

Was und Wie Sequenzieren?

- „Actionable“ Mutationen
 - Billig, klein, einfach
- Prognostisch
 - HER2 in Brust
- Prädiktiv
 - KRAS in Kolon
- Panels auch mit Markers in Trials

ABL1	EZH2	JAK3	PTEN
AKT1	FBXW7	IDH2	PTPN11
ALK	FGFR1	KDR	RB1
APC	FGFR2	KIT	RET
ATM	FGFR3	KRAS	SMAD4
BRAF	FLT3	MET	SMARCB1
CDH1	GNAS11	MLH1	SMO
CDKN2A	GNAS	MPL	SRC
CSF1R	GNAS	NOTCH1	STK11
CTNNB1	HNF1A	NPM1	TSP3
EGFR	HRAS	NRAS	VHL
ERBB2	IDH1	PDGFRA	
ERBB4	JAK2	PIK3CA	

Beadling, J Mol Diagn, 2013

