

Revisiting the Correlation of *c-kit* Mutation Status and Treatment Decisions in Canine Mast Cell Tumors

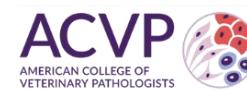
Department of Pathobiology
Institute of Immunology

Sabine E. Hammer

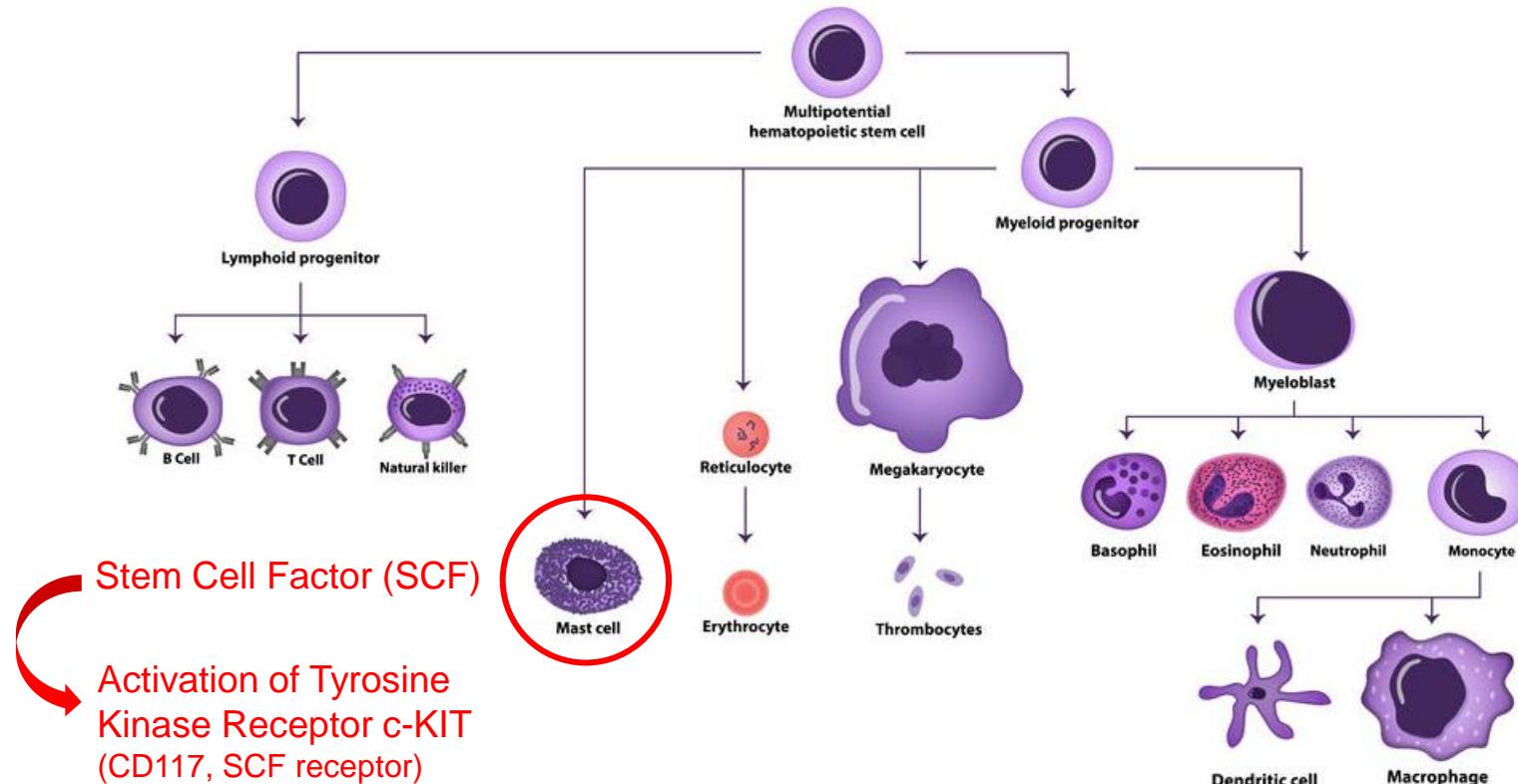
Focused Scientific Session: Diagnostic Pathology
November 14, 2022 | 8.00 – 9.45 a.m.

Relevant Financial Relationship Disclosure Statement

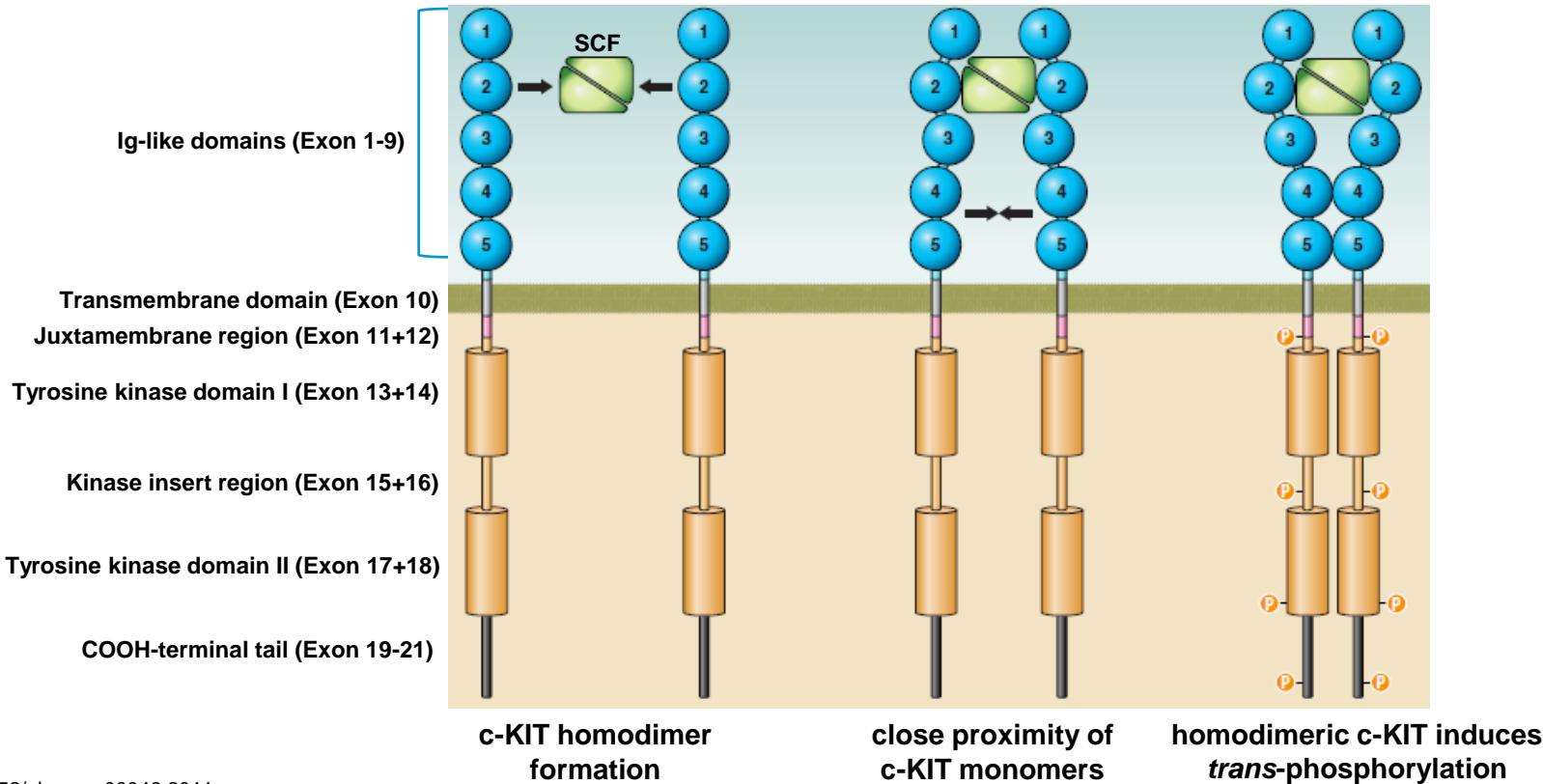
In relation to this presentation, I do not have any relationships with companies to report.



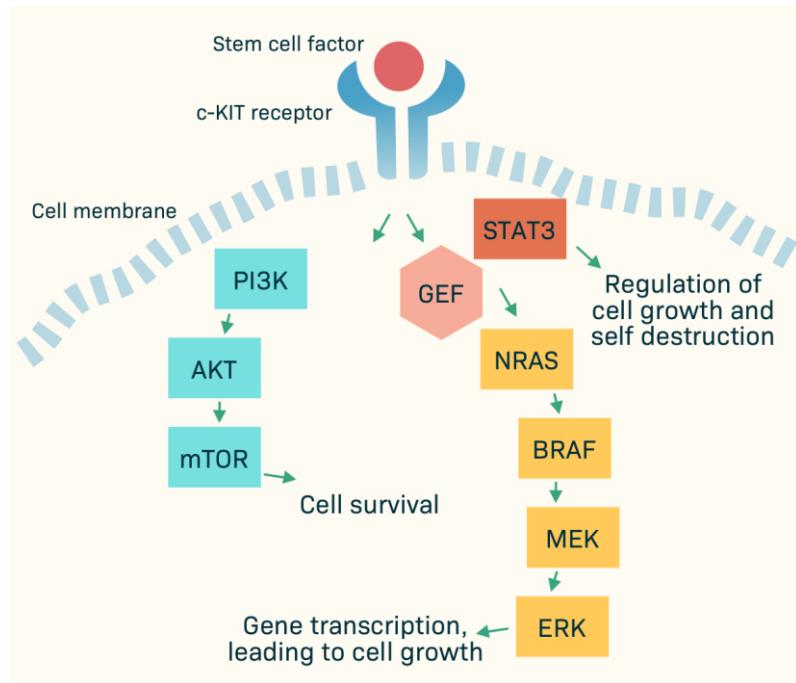
Mast Cells express Tyrosine Kinase Receptor c-KIT



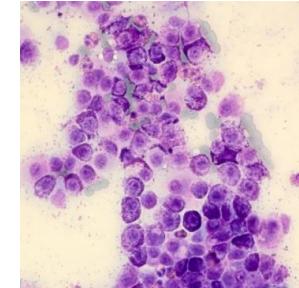
Stem cell factor (SCF) induced c-KIT activation



Stem cell factor (SCF) receptor encoded by c-kit



- Mutations in the proto-oncogene *c-kit*, encodes for the SCF receptor on the surface of mast cells → induction of constitutive receptor activation



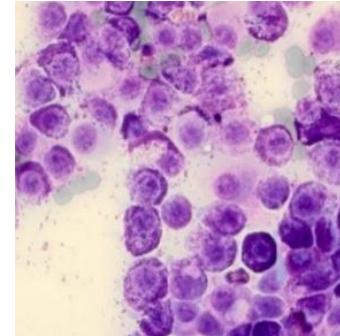
- Known mutations
 - Internal tandem duplication (ITD)
 - Single nucleotid polymorphism (SNP)
 - Insertions, Deletions
- Literature: 50% of MCTs in the dog show an ITD in Exon 8 or 11

Mast cell tumors in dog

- very common – incidence 16-21% of all skin neoplasias
- breed disposition

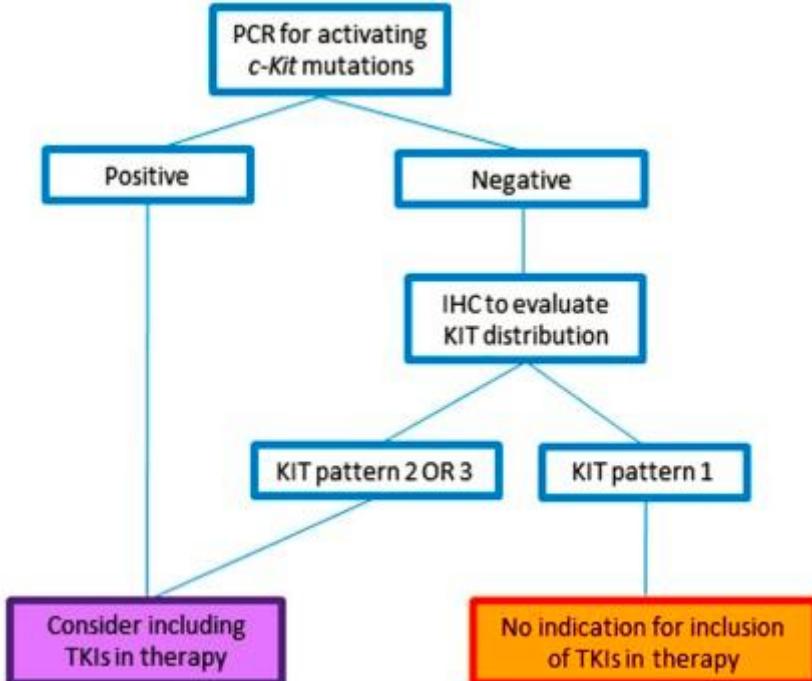


- skin solitary mass, multiple possible – "wax and wain"
- diagnosis in 92-96% of cases using cytology
no classification, no prognosis – description of malignant morphology
- classification: histopathology – Patnaik (I, II, III), Kiupel (low grade, high grade)
- therapy – surgery
recidive vs systemic disease –
chemotherapy, Tyrosine Kinase Inhibitors (TKIs)

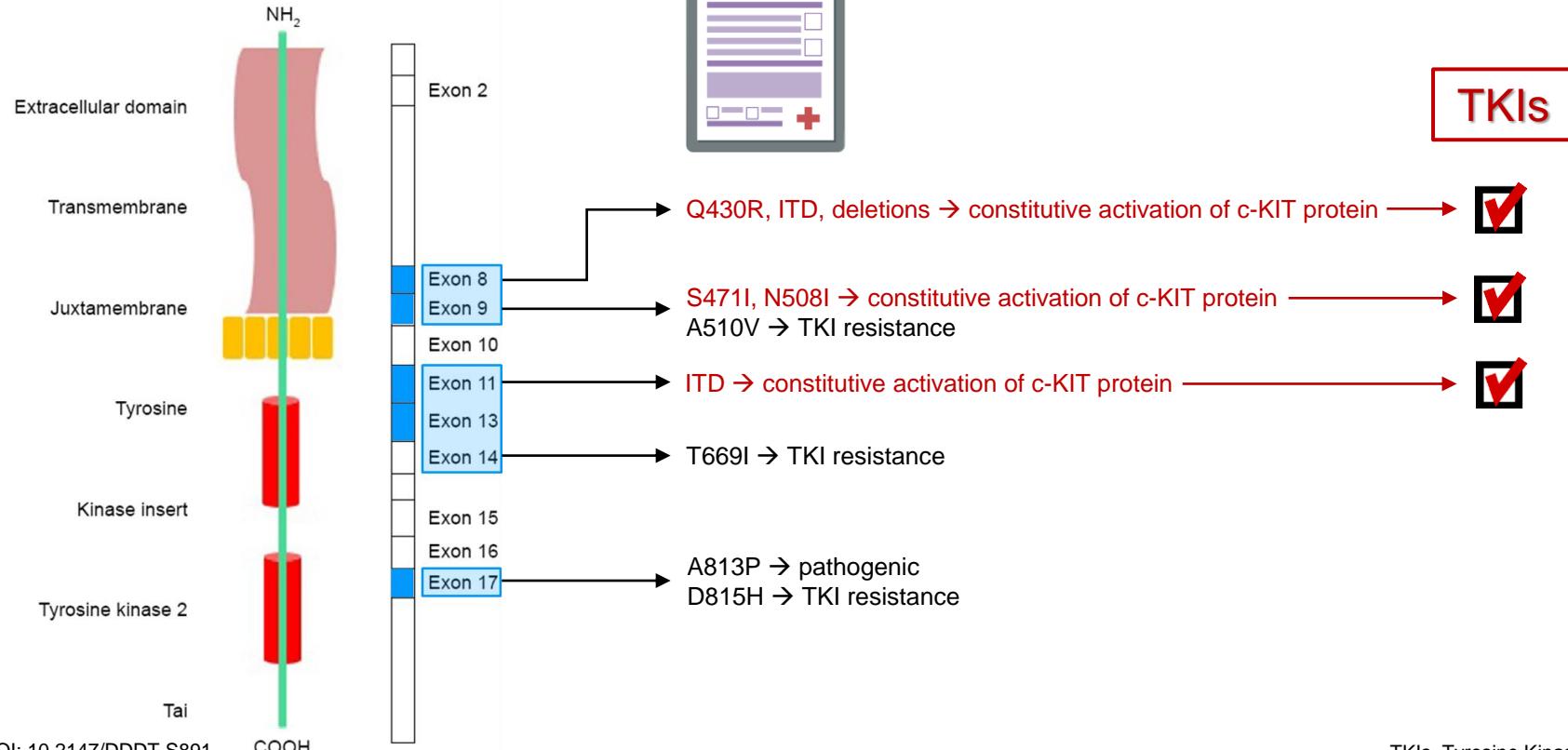


Do Tyrosine Kinase Inhibitors (TKIs) make sense?

- TKIs expand the treatment possibilities in MCT
- TKIs selectively bind to SCF receptor and inhibit its overactivation
- Is it useful to test the *c-kit* mutation status prior to MCT treatment with TKIs?
 - Yes – increased effect if activating mutation is present
 - Side effects may occur, as well as spontaneous mutations or tumor resistance

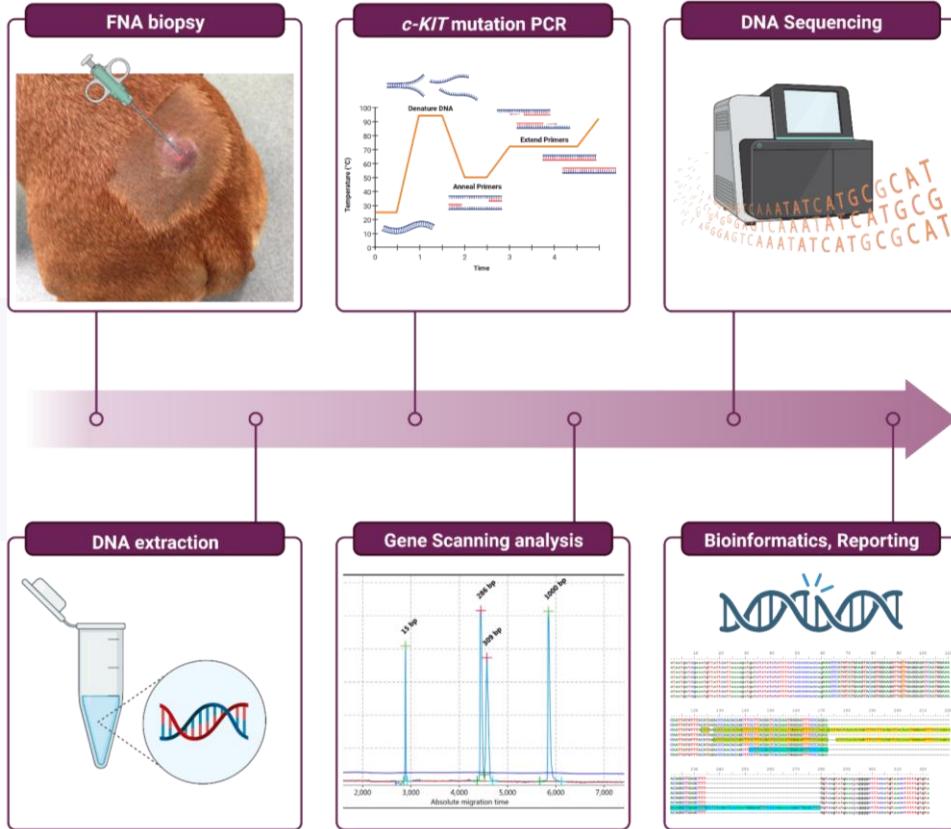


c-kit mutation screening

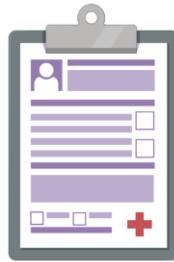


c-kit mutation analysis workflow

Created with BioRender.com



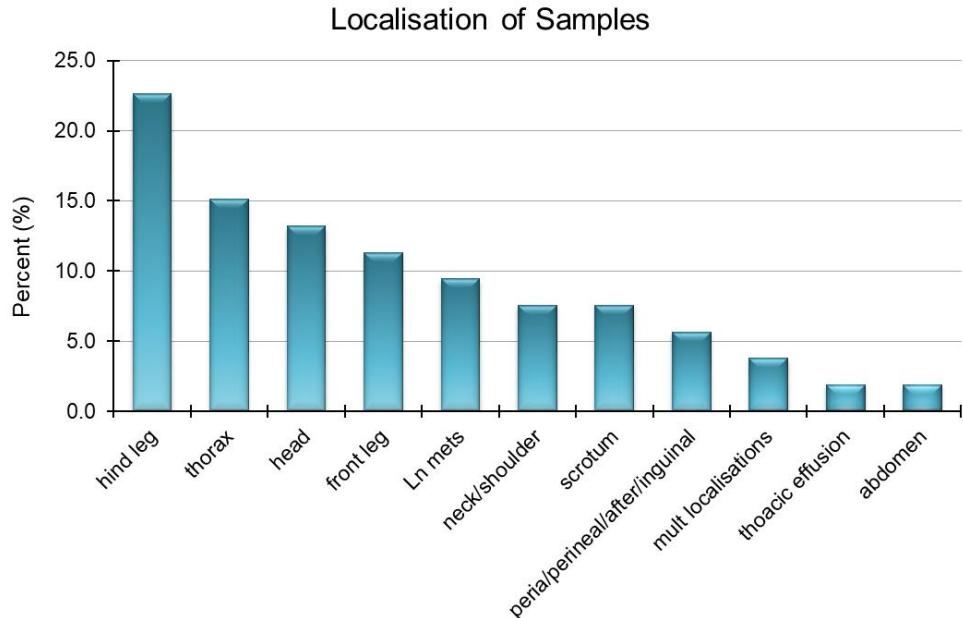
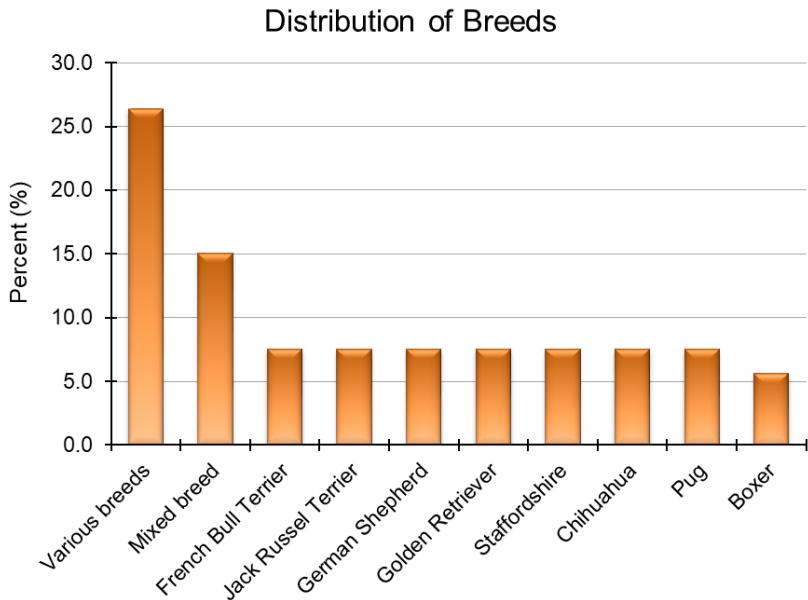
- single cell suspension (cooled or RT) as for FCM – if viable also other possibilities
- stained/unstained cytology slides – history, localization, morphology, DNA quality/amount
- histopathology samples (50µm) – history, localization, cytology report etc.



- Analysis is performed every 2nd week
- Workflow takes 4 business days
- Total turn-around-time: 14 days

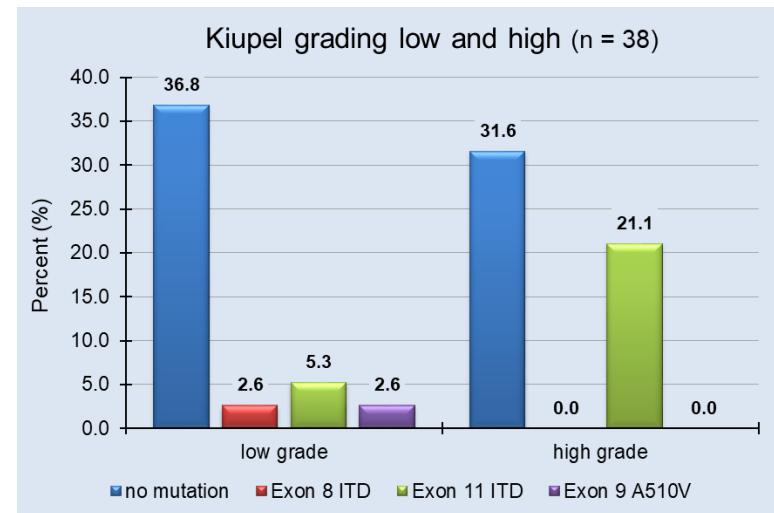
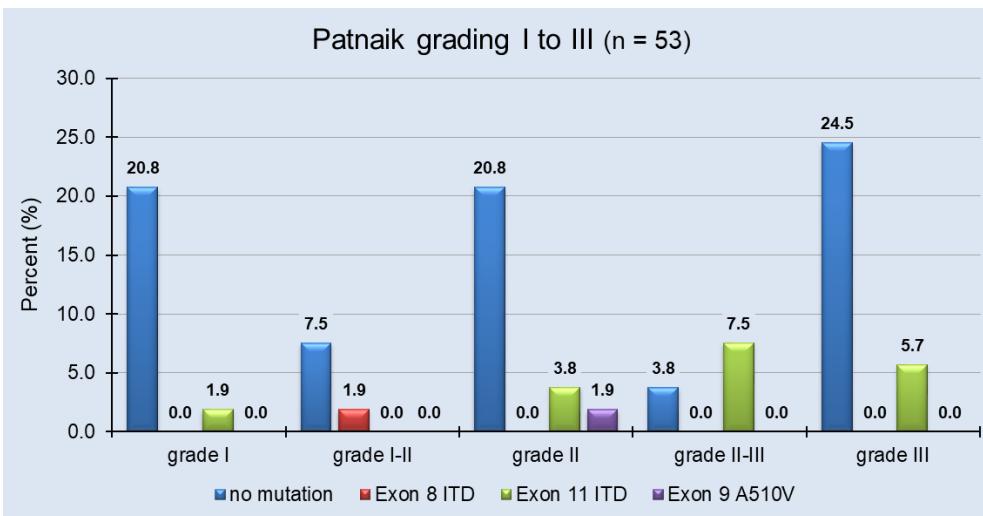
Data collected (1)

53 MCT samples - *c-kit* exons 8, 9, 11, 13, 14 and 17 analyzed



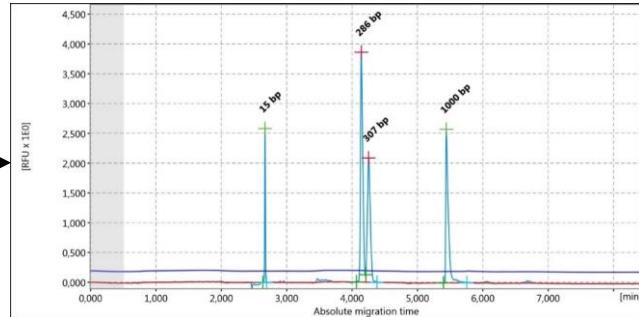
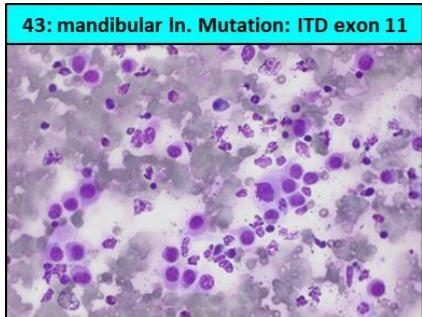
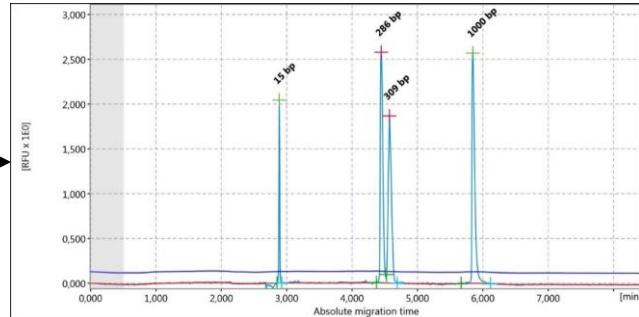
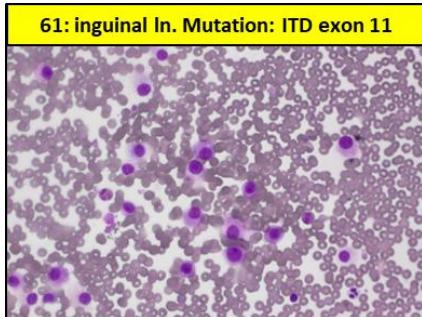
Data collected (2)

53 MCT samples - *c-kit* exons 8, 9, 11, 13, 14 and 17 analyzed
– 38 with Patnaik **and** Kiupel grading

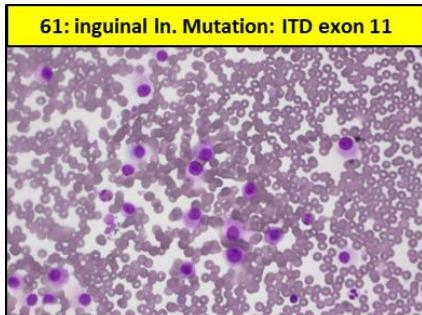


Only 20.7% patients showed an ITD in Exon 8 or 11

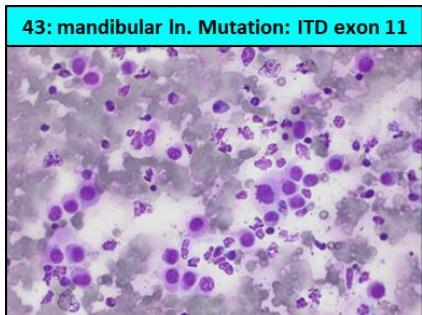
Two examples of activating mutation (ITD) in *c-kit* Exon 11 (1)



Two examples of activating mutation (ITD) in *c-kit* Exon 11 (2)



Dog Wild type	ataatgatcgaaatgttattcattaaaagatgatctrctctctttcccccacca	ggAAACCCATG	TATGAAGTACAGTGGAAGGTTGTTGAGGAGATCAATGGAAA		10	20	30	40	50	60	70	80	90	100	110
C2 Wild type	ataatgatcgaaatgttattcattaaaagatgatctrctctctttcccccacca	ggAAACCCATG	TATGAAGTACAGTGGAAGGTTGTTGAGGAGATCAATGGAAA												
C2 ITD exon 11	ataatgatcgaaatgttattcattaaaagatgatctrctctctttcccccacca	ggAAACCCATG	TATGAAGTACAGTGGAAGGTTGTTGAGGAGATCAATGGAAA												
61 Wild type	ataatgatcgaaatgttattcattaaaagatgatctrctctctttcccccacca	ggAAACCCATG	TATGAAGTACAGTGGAAGGTTGTTGAGGAGATCAATGGAAA												
61 ITD exon 11	ataatgatcgaaatgttattcattaaaagatgatctrctctctttcccccacca	ggAAACCCATG	TATGAAGTACAGTGGAAGGTTGTTGAGGAGATCAATGGAAA												
43 Wild type	ataatgatcgaaatgttattcattaaaagatgatctrctctctttcccccacca	ggAAACCCATG	TATGAAGTACAGTGGAAGGTTGTTGAGGAGATCAATGGAAA												
43 ITD exon 11	ataatgatcgaaatgttattcattaaaagatgatctrctctctttcccccacca	ggAAACCCATG	TATGAAGTACAGTGGAAGGTTGTTGAGGAGATCAATGGAAA												



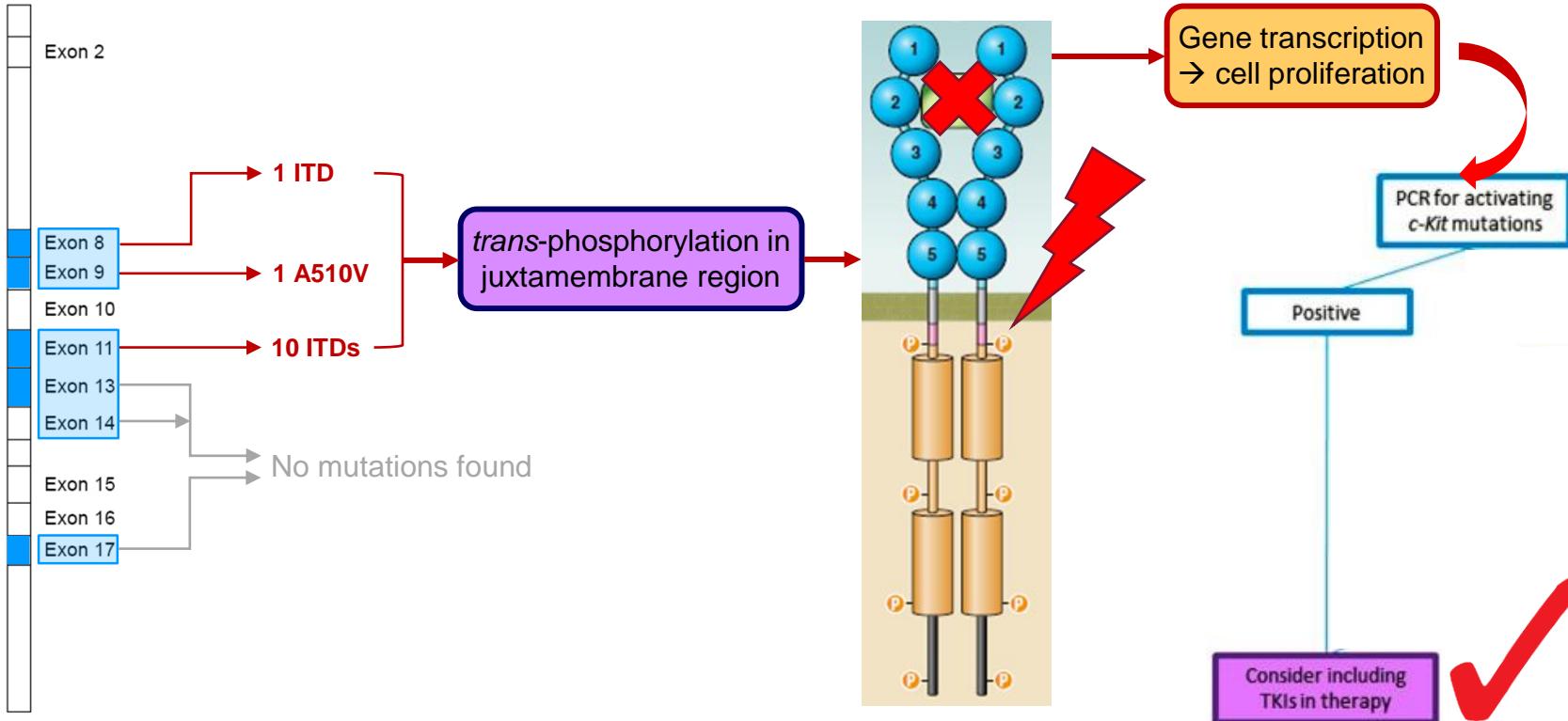
Dog Wild type	CAATTATGTTTACATAGACCCAACACAGCCTTCACGATCACAAATGGGAGTTTCCCAGAA-		120	130	140	150	160	170	180	190	200	210	220	
C2 Wild type	CAATTATGTTTACATAGACCCAACACAGCCTTCACGATCACAAATGGGAGTTTCCCAGAA-													
C2 ITD exon 11	CAATTATGTTTACATAGACCCAACACAGCCTTCACGATCACAAATGGGAGTTTCCCAGAA-	CATAACCCAACACAGCTTCCTACGATCACAAATGGGAGTTTCCCAGAA-												
61 Wild type	CAATTATGTTTACATAGACCCAACACAGCCTTCACGATCACAAATGGGAGTTTCCCAGAA-	CATAACCCAACACAGCTTCCTACGATCACAAATGGGAGTTTCCCAGAA-												
61 ITD exon 11	CAATTATGTTTACATAGACCCAACACAGCCTTCACGATCACAAATGGGAGTTTCCCAGAA-	-ACCCAACACAGCTTCCTACGATCACAAATGGGAGTTTCCCAGAA-												
43 Wild type	CAATTATGTTTACATAGACCCAACACAGCCTTCACGATCACAAATGGGAGTTTCCCAGAA-	-ACCCAACACAGCTTCCTACGATCACAAATGGGAGTTTCCCAGAA-												
43 ITD exon 11	CAATTATGTTTACATAGACCCAACACAGCCTTCACGATCACAAATGGGAGTTTCCCAGAA-	-ACAGGCTGAGCTTCTACGATCACAAATGGGAGTTTCCCAGAA-												

Dog Wild type	ACAGGCTGAGCTTT-		230	240	250	260	270	280	290	300	310	320	
C2 Wild type	ACAGGCTGAGCTTT-												
C2 ITD exon 11	ACAGGCTGAGCTTT-	Ggtcagtatgaaaayaggggcttccatgttaaccttttgtgtta											
61 Wild type	ACAGGCTGAGCTTT-	Ggtcagtatgaaaayaggggcttccatgttaaccttttgtgtta											
61 ITD exon 11	ACAGGCTGAGCTTT-	Ggtcagtatgaaaayaggggcttccatgttaaccttttgtgtta											
43 Wild type	ACAGGCTGAGCTTT-	Ggtcagtatgaaaayaggggcttccatgttaaccttttgtgtta											
43 ITD exon 11	ACAGGCTGAGCTTT-	Ggtcagtatgaaaayaggggcttccatgttaaccttttgtgtta											

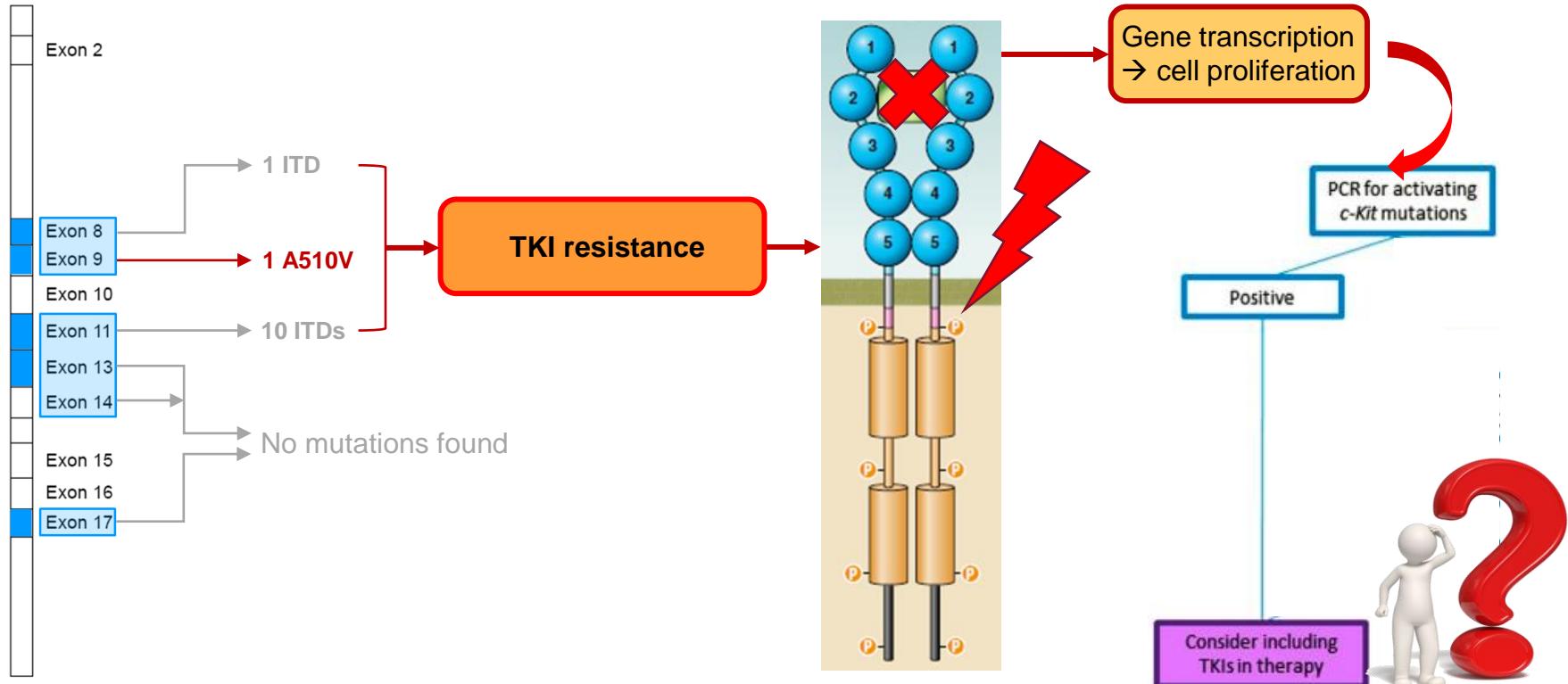
Histopathology: high-grade (Kiupel) | grade III (Patnaik)

ITD, Internal tandem duplication

Outcome *c-kit* mutation screening



Outcome *c-kit* mutation screening



Acknowledgements

**My colleagues at the Clinical Pathology, Institute of Immunology
and Institute of Pathology in Vienna
Our students Amina, Viktoria and Giovanni**



Barbara C. Rütgen

Thank you for your attention!



Sabine E. Hammer

Department of Pathobiology/Institute
of Immunology
University of Veterinary Medicine,
Vienna

+43 1 25077-2754
sabine.hammer@vetmeduni.ac.at
Veterinärplatz 1, 1210 Vienna
Austria

vetmeduni.ac.at