2023-2-3 Executive Team Meeting Notes

Date

03 Feb 2023

Attendees

Committee Member	Present	Absent
Kuffel, Gina (NIH/NCI) [C]	X	
Unknown User (leblancak)	X	
Otridge, John (NIH/NCI) [C]	Х	
Kim, Erika (NIH/NCI) [E]		Х
Sommers, Connie (NIH/NCI) [E]		Х
Debbie Knapp	Х	
Toby Hecht	Х	
Unknown User (parchmentr)		Х

Goals

• Discuss updates to ICDC and define emerging strategies and priorities

SharePoint Site

https://nih.sharepoint.com/sites/NCI-CBIIT-FNL-ICDC-ICDCLeadershipGroups

Outstanding Action Items

Agenda

Item	Talking Point	
------	---------------	--

Kuffel, Gina (NIH	ICDC Studies in the Queue
Kuffel, Gina (NIH /NCI) [C]	UBC03 Approved by SAC on 8/13/22 Transcriptomic analyses of early stage bladder cancer in Scottish Terriers detected through screening* Purdue University 19 cases Status: Submitters generating template-based data loading files OSA02 Approved by SAC on 9/21/22 "Immune Pathways and TP53 missense mutations are associated with longer survival in canine osteosarcoma* Colorado State University 108 cases Status: Submitters generating template-based data loading files ORANOIDS01 Approved by SAC on 12/5/22 "Characterization of healthy and diseased canine tissues and organoids* lowa State University Status: Submitters generating template-based data loading files UBC04 Approved by SAC on 12/22/22 "Whylole exome sequencing analysis of canine urothelial carcinomas without BRAF V595E mutation" Approved by SAC on 12/22/22 "Whylole exome sequencing analysis of canine urothelial carcinomas without BRAF V595E mutation" North Carolina State University Geases Status: Onboarding kick-off call on 1/5/23 PRECINCT01 Approved by SAC on 6/28/22 "Inhaled IL-15 Immunotherapy for Treatment of Lung Metastases" UC Davis 21 cases Status: Data not yet available PRECINCT02 Approved by SAC on 12/8/22 "Novel treatments for Diffuse large B cell lymphoma (DLBCL) in dogs" Tutts University Af cases Status: Data not yet available PRACAN01 Approved by SAC on 11/8/21 "Canine tumor mutational burden is correleated with TP53 mutation across tumor types and breeds" University of Georgia 684 cases Status: Submitters generating template-based data loading files UMCVM Approved by SAC on 4/19/21 "Comparative analysis of genome-wide DNA methylation identifies patterns that associate with conserved transcriptional programs in osteosarcoma" University of Minnesota
	44 casesStatus: On hold until FTE is designated
Kuffel, Gina (NIH /NCI) [C]	2022 BPSC Review Article Jeff Trent reached out relevant journal contacts Feedback received will require us to emphasize the connection between dogs and humans and the value of canine as a comparative model The paper has a genomic focus, however the comparative trials that exist are drug focused ldeally we can site papers and studies that highlight relevant mechanisms with a genomic facet
	Kuffel, Gina (NIH

Upcoming Steering Committee Meeting Unknown (parchme	
--	--

Minutes (Not Verbatim)

TH - Look at drug responses and the tumor itself to determine if there is a biomarker based on genomic expression. In the Intro of the BPSC paper, we could give examples of things that can be done based upon existing drug trial studies.

- AL In general, there is such a scarcity of credentialed biology in tumor progression. Drug mechanisms can be highlighted with an imaging reporter.
- DK Pre and post biopsies were attempted in Vemurafinib trial
- AL Until we have DBs to refer to and a more stable reference genome it is difficult to make progress.

Previous ICDC Use Cases from Steering Committee

- 1. Genomic correlates across platforms (DNA, RNA, protein).
- 2. Correlating multi-omics data with clinical annotation and phenotypes, particularly outcomes.
- 3. Comparative analyses of canine and human. Examples include:
 - 1. Search for conserved mutations between canine and human tumors
 - 2. Disease diagnosis (e.g. cancer type) and classification mapping between canines and humans
 - 5. Gene expression changes and mutational profiles associated with therapeutic response and outcome
 - 6. How do sporadic tumors in non-human mammals compare to sporadic human tumors?
 - 7. Correlations and model building from radiomic and pathomic features extracted from medical and histopathologic images with outcomes and genomics, as is currently being widely done with human images
 - 8. Develop biomarkers of response and resistance in humans by analyzing the responses and genomic signatures in dogs.

Previous Meeting Minutes (Not Verbatim)

- TH Circulating tumor DNA is a strong candidate for the next phase of the ICDC.
- CS There is a grant under the mammalian models from Cheryl London and co.

Action items