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Neurologic Deterioration in Children with Non-Severe Traumatic Intracranial Hemorrhage: A Multicenter Cross-Sectional Study

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OBJECTIVES/GOALS: Children with traumatic intracranial hemorrhage are monitored closely for deterioration and need for intervention. Data on risk factors for deterioration in nonsevere head injury are limited. Our objective was to identify children with hemorrhage from non-severe head injury who are at risk for deterioration. METHODS/STUDY POPULATION: We conducted a 10-site cross-sectional study of children 8. Our primary outcome was clinically important hemorrhage after injury and within 96 hours of ED arrival, defined as ED interventions (intubation, hyperosmotic agents, or neurosurgery within 4 hours of arrival) or clinically important deterioration (new or worsening signs/symptoms with an acute change in management). After testing model assumptions, we used logistic regression to identify clinical and neuroradiographic factors associated with clinically important hemorrhage. RESULTS/ ANTICIPATED RESULTS: We studied 763 children with intracranial hemorrhage, with a median (IQR) age of 3.0 (0.4, 10.5) years. Initial GCS was mild (14-15) in 89.4% (n=682) and moderate (9-13) in 10.6% (n=81). Clinically important hemorrhage was observed in 19.5% (n=149), and 7.8% (n=59) developed clinically important deterioration. Median (IQR) time to deterioration was 17.6 (4.6, 37.9) hours. In our sample, 16.3% (n=124) underwent critical interventions, 54.9% (n=419) were admitted to an ICU, and 50.1% (n=382) underwent repeat neuroimaging. We found older age (OR 1.6; 95% CI 1.3, 1.9), lower GCS (OR 5.0; 95% CI 2.9, 8.5), and epidural hemorrhage (OR 3.3; 95% CI 2.0, 5.5) was associated with clinically important hemorrhage. DISCUSSION/ SIGNIFICANCE: Clinically important hemorrhage occurred in one in five children with non-severe head injury. Clinical and neuroradiographic factors associated with ED interventions and deterioration were identified. Risk stratification algorithms using these data will be developed to assist clinicians caring for children with head injury.

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Randomized Placebo-controlled Trial to Test the Efficacy of genetically informed biomarker Nicotine Metabolite Ratio (NMR), and transdermal nicotine replacement therapy (NRT) versus varenicline Obumneke Amadi-Onuoha

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OBJECTIVES/GOALS: The overall aim of the proposed study is to evaluate the effectiveness and clinical utility of the NMR as a biomarker of response to placebo, transdermal nicotine, and varenicline to be utilized within the clinical practice as a point-of-care predictor to tailor an individual's smoking cessation treatment. METHODS/ STUDY POPULATION: A 12-week phase II, stratified multicenter, randomized, placebo-controlled trial of 3 treatment groups to measure treatment-seeking smokers (n-900:150 slow metabolizers; 150 normal metabolizers), randomized to 12-weeks of nicotine patch (active patch + placebo pill), varenicline (active pill + placebo patch), or placebo (placebo pill + patch). At the end of treatment, patients will be followed for 12 months. Study Population: Adult smokers 18–65 years old reported smoking≥10 cigarettes/day for≥6 months (verified by carbon monoxide (CO) >10 ppm). Drug and route of administration: RESULTS/ANTICIPATED RESULTS: The study would conclude the effectiveness of the interventions well defined. DISCUSSION/SIGNIFICANCE: The report will provide robust evidence to support the effectiveness of NRT intervention on smoking cessation.

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Regulation of renal function by the peroxisome proliferator-activated receptor-alpha: A novel target for treating hypertension

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OBJECTIVES/GOALS: Approximately 37 million people in the U.S. have chronic kidney disease, which is a major risk factor for cardiovascular and end stage renal diseases. PPAR-αknockout (KO) mice exhibit increased renal inflammation and blood pressure. In this study, we investigated the role of PPAR-αin renal function in a model hypertension. METHODS/STUDY mouse of POPULATION: Male 4-month-old wild type (WT) and PPAR-Î ±KO mice were instrumented with radio transmitters by artery canulation (Data Science Intl). This method minimizes stress and artifacts by avoiding the use of tethering, restraining, or anesthetizing the mice during data sampling. After recovery from surgery, we continuously measured mean arterial pressure (MAP) via radio telemetry in conscious ambulatory mice. After baseline MAP was established, vehicle (Veh; saline) or angiotensin II (Ang II) were infused using an osmotic minipump at a slow pressor dose (400 ng/kg/min) for 12 days. On day 12, we injected an intravenous bolus of fluorescin-sinistrin (3.74µl/g body weight) and collected 8 blood samples (20µl/sample) over 75 minutes to enable calculation of the glomerular filtration rate (GFR) using [GFR = $I/(A/\hat{I}\pm + B/\tilde{A}\ddot{Y})$]. RESULTS/ANTICIPATED RESULTS: Similar to our prior observations, no significant (ns) differences in baseline MAP were observed between WT and PPAR- $\hat{I}\pm KO$ mice [(mmHg): WT (n=6), 111 ± 20 vs. PPAR- $\hat{I}\pm KO$ (n=6), 113 ± 10; ns] whereas after 12 days of the slow pressor effect of Ang II, MAP was increased in both strains [(mmHg): WT (n=8), 138 \pm 11# vs. PPAR-Î \pm KO (n=8) , 156 \pm 16#; #p DISCUSSION/SIGNIFICANCE: PPAR-αprotects mice from worsening hypertension and is critical to preserving GFR during normotensive conditions. Ongoing studies are further investigating how PPAR-αregulates renal function. These finding suggest therapeutics designed to increase PPAR-αactivity could have clinical benefit in chronic kidney disease.

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The kinesin-like protein Kif11 is essential for the survival of TP53 mutant triple-negative breast cancer cells

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OBJECTIVES/GOALS: While mutant TP53 is an attractive therapeutic target in TNBC, attempts to target the mutant p53 protein directly have failed. Thus, we aim to identify pathways critical for the survival of TP53 mutant cells that can be targeted in TNBC. We have identified Kif11 as one such target and aim to further investigate its function in TP53 mutant TNBC. METHODS/STUDY POPULATION: We conducted a dual in silico/in vivo screen that identified Kif11 inhibition as preferentially inhibiting the growth of TP53 mutant TNBC. We obtained data on TP53 mutational status, KIF11 mRNA expression levels, and clinical characteristics from TCGA, METABRIC, and CCLE datasets. We treated breast cancer cell lines with the KIF11 inhibitor SB-743921. Cell counts were obtained through staining with DAPI or Hoechst and imaging on the ImageXPress PICO. We detected cell death by DRAQ7 staining and flow cytometry analysis following Annexin V-PI staining. To investigate mitotic spindle organization, we performed immunofluorescent staining with an anti-tubulin antibody and DAPI costaining. Cell cycle analysis was performed through flow cytometry. RESULTS/ANTICIPATED RESULTS: KIF11 is highly expressed in TP53 mutant and TNBC clinical samples. High KIF11 expression is associated with poorer clinical outcomes. Kif11 inhibition suppresses growth of both TP53 mutant and wild-type breast cancer cells, but preferentially induces the death of TP53 mutant cells as detected by DRAQ7 and Annexin V/PI staining. Kif11 inhibition induces a G2-M block and growth inhibition in TP53 wild-type cells. On the other hand, following treatment with the Kif11 inhibitor SB-743921, TP53 mutant cells undergo mitotic spindle dysfunction leading to the formation of multinucleated cells and cell death. DISCUSSION/ SIGNIFICANCE: These results demonstrate that Kif11 is a promising therapeutic target in aggressive, TP53 mutant TNBCs. Kif11 inhibitors, including SB-743921, have been tested in human trials, and are well tolerated, but it is unclear which patients would most benefit. Our studies show that Kif11 inhibitors may be most useful in patients with TP53 mutant TNBCs.

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The role of leucine-rich PPR motif-containing protein (LRPPRC) in myelin lipid metabolism*

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OBJECTIVES/GOALS: Leigh Syndrome, French Canadian-Type (LSFC) is a neurometabolic disorder caused by mutation of mitochondria-related gene, LRPPRC. White matter lesions and demyelination in central nervous system are common in LSFC. LRPPRC is enriched in myelinating glial cells, yet its role is not known. Our goal is to elucidate its mechanistic role in myelination. METHODS/STUDY POPULATION: We crossed C57BL/6N mice bearing a LRPPRCloxP allele with mice bearing a Plp-CreERT2 allele. Mice with the Plp-CreERT2 allele expresses a tamoxifen-inducible Cre under the control of the Plp promoter, which drives expression in oligodendrocytes. Using these strains, we can target the deletion of LRPPRC, via tamoxifen injection, in both newly formed myelin and mature myelin. Plp-CreERT2; LRPPRCL/L (LRPPRC-KO) or control littermate mice will be injected for LRPPRC deletion at developmental and maturation stages of myelin. Immunofluorescence and electron microscopy of isolated brain tissues will be used for myelin integrity analysis. Cognitive functions of the mice will be measured via behavioral tests. Lastly, we will submit tissues for lipidomic analyses to observe any lipid metabolite variation. RESULTS/ANTICIPATED RESULTS: Behavioral and motor defects would be expected in LRPPRC-KO mice performing in cognitive function tasks across myelin maturation stages. Electron microscopy-based structure analysis of optic nerve, corpus callosum, and spinal cord should reveal thin or loss of myelin on the axons of LRPPRC-KO compared to control. Immunofluorescence staining of major myelin structural proteins, including myelin proteolipid protein (PLP), myelin basic protein (MBP), and myelin-associated glycoprotein (MAG) would be expected have lower levels in LRPPRC deficient tissues. Since myelin is a lipid-rich species, we would also expect lipid concentrations to be affected. LRPPRC-KO lipidomic analyses of myelin-related lipids should depict lower levels in comparison to control, which would imply dysfunctional lipid metabolism. DISCUSSION/SIGNIFICANCE: There are limited studies in ameliorating neural deficits caused by LS and LSFC. Successful completion of this project would help elucidate the functions of LRPPRC in myelination and lipid metabolism and potentially provide insights for developing novel therapeutic strategies for alleviating the demyelination and neural deficits in LSFC.

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The Role of the IL-6-IGF-II Axis in Systemic Sclerosis-Associated Lung Fibrosis

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OBJECTIVES/GOALS: Interleukin (IL)-6 is produced in excess in Systemic Sclerosis (SSc). Likewise, microarray analysis of Insulin-like Growth Factor (IGF)-II-treated NL fibroblasts revealed increased expression of the basic helix-loop-helix transcription factor, BHLHB2. Our goal is to delineate the role of BHLHB2 in the fibrotic response to IGF-II and IL-6. METHODS/STUDY POPULATION: Primary lung fibroblasts were cultured from human lung tissues at 37oC and 5% CO2. Cell cultures were stimulated with IL-6. Gene expression was measured using quantitative PCR (qPCR). IGF-II mRNA expression levels after IL-6 stimulation were compared with those of the housekeeping gene PPIB (Peptidylprolyl Isomerase B). Western blot was performed on nuclear and chromatin-bound subcellular fractions from treated lung fibroblasts. BHLHB2 protein levels were assayed in response to IGF-II in comparison to PBS as vehicle control. RESULTS/ANTICIPATED RESULTS: Results: Our results show that IL-6 increases IGF-II levels in fibroblasts. In turn, IGF-II increases BHLHB2 nuclear localization. We further show that IL-6 increases BHLHB2 levels and its nuclear localization in lung fibroblasts. Our findings are novel since the role of the transcription factor BHLHB2 in the IL-6 induced/ IGF-IImediated fibrotic response in SSc lung disease remains unexplored. DISCUSSION/SIGNIFICANCE: Our findings may provide a rationale for combination therapy to block IL-6 and IGF-II function concomitantly and thus halt the progression of SSc pulmonary fibrosis (PF). Our findings may have wide implications for lung fibrosis associated with various diseases, since SSc-PF, is characterized by the activation of common fibrotic pathways.

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A Pilot Study of Tear Cytokine Profiling in a Patient with Ocular Graft versus Host Disease

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OBJECTIVES/GOALS: Ocular graft versus host disease (oGVHD) affects ~50% of individuals after an allogeneic hematopoietic stem cell transplant for treating blood cancers. OGVHD results in severe