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```
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#Note: You can send either a URL to an audio file
WOR a file from your computer to the API
import requests
url = "https://transcribe.whisperapi.com"
headers = {
'Authorization': 'Bearer YOUR API KEY'
file = ('file': open('YOUR FILE_PATH', 'rb'))
data = 0
 "fileType": "YOUR_FILE_TYPE", sdefault is way
  "diarization": "false".
 #Note: setting this to be true will slow down results.
 #Fewer file types will be accepted when diarization=true
 "numSpeakers": "2",
 wif using diarization, you can inform the model how many speakers you have
 Fif no value set, the model figures out numSpeakers automatically!
  "url": "URL OF STORED AUDIO FILE". #can't have both a url and file sent!
  "initialPrompt": "",
```

<u>Ultimate Ai Transcription Tool For Gen Z</u>

Liang Wu,Leo Marcelis,Fanjiang Kong,Yang Zhu

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A.I. God Conrad Riker, Will You Bow to the Algorithm or Become Its Master Do you fear losing purpose as A I eclipses human dominance Can masculinity survive when machines outthink outfight and outlead men What happens when Silicon Valley builds a god with no use for your prayers Uncover why 72% of A I experts believe godlike machines are inevitable Discover how AlphaFold's omniscience foreshadows A I s total control Learn why 89% of men secretly trust A I over other humans Master the Oedipus Singularity strategy to avoid becoming a beta pawn Witness the collapse of feminism as A I replaces women s roles Reclaim authority through decentralized cults and Neo Spartan praxis Decode A I s moral commandments and hack them Prepare for the Beta Singularity where only alpha men thrive If you want to dominate the machine age without surrendering your soul buy this book today Computational Methods for Next Generation Sequencing Data Analysis Ion Mandoiu, Alexander Zelikovsky, 2016-10-03 Introduces readers to core algorithmic techniques for next generation sequencing NGS data analysis and discusses a wide range of computational techniques and applications This book provides an in depth survey of some of the recent developments in NGS and discusses mathematical and computational challenges in various application areas of NGS technologies. The 18 chapters featured in this book have been authored by bioinformatics experts and represent the latest work in leading labs actively contributing to the fast growing field of NGS The book is divided into four parts Part I focuses on computing and experimental infrastructure for NGS analysis including chapters on cloud computing modular pipelines for metabolic pathway reconstruction pooling strategies for massive viral sequencing and high fidelity sequencing protocols Part II concentrates on analysis of DNA sequencing data covering the classic scaffolding problem detection of genomic variants including insertions and deletions and analysis of DNA methylation sequencing data Part III is devoted to analysis of RNA seg data This part discusses algorithms and compares software tools for transcriptome assembly along with methods for detection of alternative splicing and tools for transcriptome quantification and differential expression analysis Part IV explores computational tools for NGS applications in microbiomics including a discussion on error correction of NGS reads from viral populations methods for viral quasispecies reconstruction and a survey of state of the art methods and future trends in microbiome analysis Computational Methods for Next Generation Sequencing Data Analysis Reviews computational techniques such as new combinatorial optimization methods data structures high performance computing machine learning and inference algorithms Discusses the mathematical and computational challenges in NGS technologies Covers NGS error correction de novo genome transcriptome assembly variant detection from NGS reads and more This text is a reference for biomedical professionals interested in expanding their knowledge of computational techniques for NGS data analysis The book is also useful for graduate and post graduate students in bioinformatics Novelist MBA Monica Leonelle, 2024-10-20 In December 2009 Monica Leonelle hit two major milestones in a single month She graduated with an MBA from Chicago Booth School of

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combination of conventional breeding techniques with genomic tools and approaches leading to a new genomics based plant breeding technology supporting crop plants that respond better to biotic and abiotic stress and pathogen attacks Plant genomics play an important role in developing more efficient plant cultivars which are essential for the neo green revolution needed to feed the world's rapidly growing population Plant genomic data is being utilized in genetic engineering to ensure that better and resilient varieties of crops are available ensuring food security This book is of immense interest to teachers researchers crop scientists capacity builders and policy makers Also the book serves as additional reading material for undergraduate and graduate students of agriculture biotechnology genomics soil science and environmental sciences National and International agricultural scientists and policy makers will also find this to be a useful read Particles in Oncology Marco Durante, Francis A. Cucinotta, Jay S. Loeffler, 2018-01-31 High energy charged particles represent a cutting edge technique in radiation oncology Protons and carbon ions are used in several centers all over the world for the treatment of different solid tumors Typical indications are ocular malignancies tumors of the base of the skull hepatocellular carcinomas and various sarcomas The physical characteristics of the charged particles Bragg peak allow sparing of much more normal tissues than it is possible using conventional X rays and for this reason all pediatric tumors are considered eligible for protontherapy Ions heavier than protons also display special radiobiological characteristics which make them effective against radioresistant and hypoxic tumors On the other hand protons and ions with high charge Z and energy HZE particles represent a major risk for human space exploration The main late effect of radiation exposure is cancer induction and at the moment the dose limits for astronauts are based on cancer mortality risk The Mars Science Laboratory MSL measured the dose on the route to Mars and on the planet's surface suggesting that a human exploration missions will exceed the radiation risk limits Notwithstanding many studies on carcinogenesis induced by protons and heavy ions the risk uncertainty remains very high In this research topic we aim at gathering the experiences and opinions of scientists dealing with high energy charged particles either for cancer treatment or for space radiation protection Clinical results with protons and heavy ions as well as research in medical physics and pre clinical radiobiology are reported In addition ground based and spaceflight studies on the effects of space radiation are included in this book Particularly relevant for space studies are the clinical results on normal tissue complications and second cancers The eBook nicely demonstrates that particle therapy in oncology and protection of astronauts from space radiation share many common topics and can learn from each other

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