we are honored to have dr jerry collins dr collins associate director of the developmental therapeutics program in the division of cancer treatment and diagnosis at the nci he received his phd in 9 from the university of pennsylvania and completed a postdoctoral fellowship in clinical pharmacology at johns hopkins he has authored or coauthored over 00 papers in the field of clinical pharmacology primary emphasis in the area of pharmacokinetics and pharmacodynamic principles in the field of cancer prior to his current position he spent years at the fda please enjoy todays lecture my name is jerry collins im here today to talk about topics that are contained in chapters and of the textbook my day job is that i lead the developmental therapeutics program at the national cancer institute

my background is that ive spent half of
my adult life at the national cancer
institute and half of my adult life at
the food and drug administration
neither of whom assumes responsibility
for anything that i might say tonight

chapter was

written by bob dedrick who originally
joined me in sharing this but hes
retired left me his slides and ill have
to try to represent them as i can bob

dedrick

was a chemical engineer he was a founding director of the chemical engineering section here at nih and just had a marvelous perspective about the relationships between phenomena in

the physical chemical world
and phenomena in the biology world
so what do chemical engineers do well
one of the major things they do is they
make enormous quantities
of materials specialty chemicals bulk
chemicals kilograms in these large

chemical plants

and how do they do that

well they start out figuring out how to synthesize the molecule in test tubes using milligram quantities and so the process of going from milligrams to kilograms which is at least a million fold difference is what chemical engineers call scale up

and as bob thought about that process his his lifelong training his question is well whats it like in biology

in biology we have these just within mammals we have these tiny rodents and we have these enormous animals what are the similarities and differences how do you get from small creatures to large creatures what are the similarities and what are the differences in fact one of bobs favorite sayings is

biologists

engineers look for similarities

look for differences and most of the people who signed up for this course are are of course biologists but

well try to give bob a chance to uh

convince us of the role of

similarities so this is a lab tech in

building and campus a little bit

earlier in time when you didnt have to

wear gloves all the time when you were

handling animals and so the mouse and

the rat clearly

have similarities you can see that they

look

sort of the same shape and different from other mammals but its a tenfold difference youre talking about a 0 gram mouse and a 0 gram rat so what are the implications of that size other other than uh one needs more food to eat

and

you know more cage space

but bob didnt stop it there his other

hobby was that he was a photographer

so

sure enough he wandered down to the
washington zoo and took this picture
so the picture he convinced the
zookeeper who was happy to participate

in this experiment to stand next to the elephants that the zoo keeper worked

with

and so compared to a rat a human is 00 times larger

but

compared to

a human

a mouse is several hundred times larger than a human at least in terms of body weight so we just have this enormous

scale

just again just within the mammalian kingdom in terms of um what size processes were bobs a very curious

person

he knew how that worked in uh chemical engineering and he wanted to learn and

contribute to that in

the world of drug distribution and

pharmacokinetics

so it turns out that

because theres such a wide variation in body weight that biological processes

have to scale

they they do things differently they do

things at different rates at different

turnover

volumes at the small end of the scale than the large end of the scale

and

0 years ago

adolf published this paper in science in

9

on allometry

and everybodys eyes glazed over and said what why do i need to know anything about elementary well adolfs simple

point was

even if you dont know

the fundamental underlying processes

that youre measuring whatever it is

temperature heart rate heart rate

amount of food intake

body surface area

theres an empiric relationship that

covers

all of those

diverse equations and thats this

equation in the middle of the slide

where the property

is equal to some constant times body

weight to the nth power

well thats all that it means it doesnt

mean that we know exactly why

a is one value and m is another value it

just means that as you as you look at

series of data across a wide range this

equation comes up over and over again

and for someone whos curious they would

say this cant be a coincidence in fact

this is very similar to the way chemical

plants are designed theres a scale up

from milligrams to kilograms that

follows a uh allometric allometriclike

like process

when you see these plots theyre a
little different from most of the
pharmacokinetic plots youve seen in
this course in that its a log log plot
both the y and the xaxis are logarithms
its the only way you can squeeze all

a diverse mammalian species onto the

the data from

same graph

and this the two squiggly lines are just showing you what happens when the property is directly proportional to

body weight where the exponent is

versus when its 0 which

is sort of a magic number in a llama

tree

good news is we wont talk about why
its a magic number so here here is uh
heat production

so

all mammalian species are warmblooded animals theyre aiming for a particular target

that they achieve that

primarily except for us they achieve it

primarily by adjusting their rate of

metabolism adjusting their their heat

output

and if you plot on the bottom everything
from a mouse to an elephant over you
count them half a dozen logs
and then you look at the heat production
on the yaxis

it it keeps coming up again no matter
what property you are looking at
um the allometric equation the loglog
plot is important

so this is the way bobs thought process
works he says well there must be a lot
of similarities there must be a way we
can put this together in a way that
would be useful in drug distribution and
pharmacokinetics

so

more recently a mere 0 years ago bob
and his colleague collaborator ken
bischoff invented something called
physiologic based pharmacokinetics
now largely pharmacokinetics today

empiric based on one compartment and two compartments

remains

of questions but there are a number of questions that are much better answered or answered in more detail if you think about the underlying biology the underlying physiology so lets go through a couple of answers one of the one of the reasons i like

bobs uh

publication on cytoscene arabinoside rsc is that he scaled the size of the boxes to represent the volumes of the

compartments

and most of the empiric models that i see all the boxes are the same size

thats okay

but it sort of gives you an idea of how important the muscle mass is we rarely think about muscle mass in terms of the implications of pharmacokinetics and pharmacodynamics but its the largest reservoir in many cases of where the

ah the part youve been waiting for a mass balance equation particularly a differential equation

drug goes

all thats meant by a mass balance equation is take a look at an organ or an organism

figure out everything that goes in and comes out think about whats happening in the end and write it down and see if you learn anything about the processes and about that organ and about that organism so what are those things for metabolism theyre things like vmax and km for transport that starts with just

various flow rates q and then for other
elimination processes we talk a lot
about clearance cl
lets see what we learned from that so
this is one of the this is is the model
that ive been talking about bobs model
in uh for heresy
and this is his simulation

for

area c data at the bottom the bottom
line and the sum of are c plus area u
its metabolite at the top which is
important because the assay a lot of
people used in those days was total
radioactivity which is mostly measuring
the metabolite
so do those lines fit those data points
i think everybody in this room could
probably do a better job
fitting those data points to a
curvilinear line but bob didnt fit

curves

those he predicted those

he simulated the behavior based on all those micro constants vmax km blood flow

and total clearance and then taken in that context its pretty remarkable that you get anywhere near

the

empiric data on pharmacokinetics
thats one species
one drug does that generalize again
bobs interested in principles that you
know go across a a wide landscape
so this is another cancer drug um

five flurry uracil

and

over the years data were collected in uh
dogs humans rat and mouse
and they all look a bit different
uh when plotted when coplotted on the
same graph where all youve done is
normalize the yaxis by uh by the dose
wonder why that is i wonder what whats
happening there can the same structural
pharmacokinetic model
predict the behavior for four different
species

if you know the differences in the in the micrograde constants and sure enough you can do that but a

more important or more interesting

question arose is

you know its not just size that changes

across mammalian kingdom its also other

properties like time

heart rates uh are radically different

across uh

species

lifespan is radically different and it

seems like

each species has its own time clock

and can you correct for that in trying

to understand differences across species

well

bob is far too humble to have called

this a dedric plot but

core and colleagues published this in

the

late 90s in which they named it after

bob in which all they did

was scale the time axis by body weight

and the concentrations all fall on the

same line

okay

in the grand scheme of things what does

that mean

again the fundamental message over and over again is

you can find similarities even when there appears to be a lot of differences how hard is it to get those data to do

those simulations

well in the metabolism lectures uh in back in the fall you heard some talk about doing hepatocytes and microsomes

and various ways of

looking at metabolism

of drugs in the body so this is a set of i dont know drugs that was published

by

brian houston

in

in the earlier in the 90s in which he looked at hepatocyte cultures okay you could you can grow hepatocytes in culture and this happens to be rats so were switching from human which was area c to rat now and instead of looking one drug were looking at different drugs and what brian houston and his colleagues did was create a physiological pharmacokinetic model

which related

in vitro benchtop measurements and hepatocyte systems to what that would look in the whole body of a rat and then in their laboratory they also collected the whole body data pretty remarkable curve if you look at any of those points theres deviations but and theyre sometimes theyre the most important story but the key is were looking over three orders of magnitude four orders of magnitude and were seeing similarities across a wide range despite the fact that there are radically different structures that are are the drugs that

were

studied here

a few years later did the same
experiment that the houston lab did
um except they looked at humans and they
looked at a few more drugs in i
didnt count them but maybe theres 0
of these
the interesting thing is that that

straight line falls apart as you get

very low

and as you get very high it looks very similar why is that

well

if youre interested in similarities you focus on the right half if youre interested in differences you folk on the left both are important if we go

back

to what what was the fundamental rate of clearance in the rat the curve ends at 0

in terms of clearance units on the

yaxis

whereas

in humans

its tails off at 0 what does that
mean there are other things in the body
besides the liver that account for how a
drug is biotransformed and excreted from
from the from the body so in general you
would not expect metabolism by the liver

alone to do it

it works in the rat because the rat is so incredibly efficient at transforming drugs relative to humans

so when humans

have drugs that have low clearances then you begin to see to parse out this tail on the left side of the graph which shows the nonhepatic effect so again you learn something from similarities you learn something from differences the last thing i i want to say of course

ive already said that

engineers look for similarities

biologists look for

differences

but both have in common

the fact that they collect a lot of data
in the laboratory and in other systems
and they want to use those data to
predict or extrapolate or scale up
from micro or many systems to larger
scale apparatus or larger scale animals

and so

always the engineer bob says that

biologists are

have a strong

interest in similarity just like

engineers do

all right lets transition away so so in

summary bobs idea and approach this is
to relate the things that he learned in
the physical chemical world the things
that are important in the biological
world particularly for uh for drug drug
development

and that was very helpful in stimulating such areas as physiologic based pharmacokinetics

which

took 0 years before about 0 years ago
it started to go into the mainstream in
in clinical pharmacokinetics so some
ideas take longer than others
what are we doing here at nih well
on about that same time frame about 0
years ago

nih started to think about reinventing portions of it and the nih roadmap was one of the mechanisms for doing that and one of the priorities in terms of rethinking some of the missions of nih was reengineering the clinical research

nih is always and forever will be known for its basic research thats thats the

enterprise

strength but since the basic research is
supposed to lead to some
payback later on in terms of the
clinical domain perhaps a little bit of
thinking in that area is is would be

useful

one of the things that came out of that

process was a creation of a new
institute or center the national center
for advancing translational sciences
whose mission statement says
they would like to speed the delivery of
drugs diagnostics and devices to
patients

its not that the folks that and cats
are in building one at nih think that
basic research is no longer important it
just means that we need to pay a little
bit more attention and we need to have a

critical mass

in terms of

the practical aspects of drugs

diagnostics and

devices

across the nih campus in various

institutes

the institutes have had small
therapeutics development or diagnostics
development programs for a long time
most of them dont have a critical mass
but they they they do perform a role
within within those institutes so it

wasnt

a completely out of the box idea it was just more like shining the spotlight on

it

now

in the program that i lead developmental therapeutics we do this with a much

larger

mission so

lets do the math 0 years ago congress passed a law that mandated nci

to

new therapeutics for cancer
if you think the landscape is difficult
for cancer therapeutics today 0 years
ago it was even more miserable
and congress said lets do something
about it there may be a parallel system
going on in the current political

climate but

were government employees and dont
talk about politics in any event our
website dtpcancergov
is the program that i lead and our major
pipeline is called nextcancergov
and the commercial
so chapter is about first in human
studies which is an area that i spent
most of most of my life most of my
professionalized working on
i am also like bob a chemical engineer
by training but went into a postdoctoral
fellowship in clinical pharmacology
showed up here at the nih and um

i

about the interface between

pre clinical and nonclinical studies

and

by and large design

pivotal phase trials they do a far

better job than i ever would but im

interested in that interface between

the last of the preclinical studies and

the early clinical studies so over the
last two weeks ed sauceville and chris
takimoto have been trying to line you up
to think about the transition between
the late stages the transition between
the late stages of preclinical
development and how that influences the
earliest earliest clinical phases and
you can find in any textbook or review
article a diagram like this
and it is a nice model system to think
about but it certainly doesnt reflect
the way all trials are done increasingly
today

folks will say ive seen some really exciting results in a tiny number of patients in first in

human phase studies

im going to make a large gamble and go

to an incredibly expensive large

complicated phase study

thats a big gamble

the success rate is relatively low but

when it turns out you gain years almost

a decade in some cases

and you advance an effective therapy

into the clinic

but its a big gamble when you go outside this paradigm similarly

drug gets

perspective gets to phase two and it
looks like it has this little problem
that it doesnt have much activity
and so theres a panic
and you certainly dont want to go
directly to phase three just because you
finished your phase two
if the result of your phase two is you
didnt find much activity so you throw
it back to the lab and you say fix this
get me a similar molecule that actually
has that activity so this is the real
world theres a lot of back and forth a

to the clinic

lot of iterative processes until you get

this slide just talks about the

contemporary trend to reengineering

first in human studies theres a lot of

new players cancer is a more attractive

target for many drug development

sponsors commercial and noncommercial

and just many things are coming together
to challenge the traditional way that
weve developed drugs are there some
better ways

well

i cant count the number of different
ways that you can do a first in human
study there are many of them
and my advice is not to get too attached
to any of them

this is an era in which theres enormous
flux in the philosophical underpinnings
of what a first in human study is is all
about and there are radical changes that

occur all the time

i would encourage you to think about the fundamental principles that no matter what algorithm you use for conducting a

first in human study

you have to do the scariest thing in

drug development

you have to pick the size of the first

dose in the first human being okay so

all the modeling all the preclinical

studies

all the genomics in the world

dont you know help help reassure you that you have the right starting dose

but

you generally use a pretty large safety
factor you calculate what you think the
right dose will be in humans
and you put in a safety factor

and

you hope everything goes well if
everything goes well your reward is you
have a second problem
where do you go from the first dose the
first dose is rarely ever the right dose
the right dose in either one that does
it is basically the therapeutic index it
either doesnt have enough activity or
it has too much toxicity

so youre going to have to go somewhere up or down usually because youve had a large safety factor youre going to have

to escalate

that is almost as scary
as picking the first dose why is that
because if you escalate too quickly and
you have a steep slope a narrow
therapeutic index

you can go from a dose thats pretty well tolerated to one thats

lifethreatening

pretty scary i mean underneath

everything we

first do no harm but its a tough balance between doing no harm and being

timid

whats the downside of being timid well if youre an investigator if youre

a patient

the downside is youll get homeopathic

doses

youll get doses that have no chance of improving your treatment at all

so

in order to maximize the potential benefit to patients you have to think of

ways of

doing escalations that we never considered before like escalating within

the same patient

by taking larger steps

um

if youre an investigator at the time that i first came into this

field thered just been a series of
first and human studies at large cancer
centers that took three years to
complete

so from the time the first dose was safely given

until the time that the final dose
which is just the beginning of the
process of figuring out whether theres

any activity was defined
that creates a terrible discontinuity
between among the clinical studies
clinical fellows who are designing this
study the clinical fellows who are
running the study and the clinical
fellows who are interpreting it and
trying to write it up afterwards

so its

efficiency is not just a fancy word
efficiency is very important for
patients to get
potentially exciting therapy for

to move along with their career and to keep the whole thing going the conflict

is between

investigators to

caution

safety efficiency and efficacy no matter
what design you choose to use
youre still going to have to face that
that factor

for um

0 years there was something called the modified fibonacci escalation scheme what that meant was that when you submitted a protocol to your irb or ethics committee the escalation procedure was predefined and that you would follow this mathematical algorithm for for increasing the doses

so your safety effect if you look at the low end of the scale 0 if you have a 0 safety factor

then your starting dose is 0

and if thats safe

where do you go next well in general in

balancing

safety versus a desire to get some

efficacy

you assume i had such a big safety factor i want something big for my first

step so you double it

all right

and everything is still okay thats the

usual outcome

but youre nervous about continuing the

doubling because

you dont know what you dont know you

dont know where you are in humans

nobody else has been there before

and so this idea that you would

serially reduce the incremental step in

terms of percent escalation

very roughly tied to the fibonacci

series in mathematics became the gold

standard for how

first in human studies were done

again

many inefficient studies because you

reach you cone down to such a small step

that youre

youre youre working hard against inter

subject variability

if you have a dose escalation step of 0

percent

and intersubject variability of 0

you have to be careful about the noise

so a number of adaptive uh designs have

been proposed

not just by myself and my colleagues bob

dedrick bruce chavner at al

but by others in which they try to think

of ways of

finetuning the escalation

scheme so that its no longer predefined

so you tell the irb what your starting

dose is and you tell them how you will

make your adjustments but you dont lock

it down in concrete

so lets talk about what those

principles might be

well

this is a pkpd course

so our fundamental thought

is that concentration circulating in the

body

tells you at least a lot not everything

but tells you a lot about whats likely

to happen in terms of toxicity and

efficacy

indeed

the concentration of drug circulating in

the body

is considered a biomarker
or an end point in many clinical studies
we expect in the toxicity domain
equal toxicity for equal exposure first

is all about hoping you have a terrifically effective drug

in human

but

the first principle is is it safe
can this structure be given safely to
human beings

so how might you do an adaptive design well pay attention to what you already know so if you um give the biggest dose you think you can to humans you compare it to to start with you compare it to how large are those animals tolerated all right so thats just on the basis of those and you know thats one way to escalate but when you consider interspecies differences in metabolism youre probably much more interested in blood levels or concentrations in plasma because thats what pharmacodynamic effects are are built upon so you can change your escalation strategy to a

from a predefined set of numbers
on a universal scale to strategy that
says the size of my escalation steps
will be determined by whether it appears
that were very far away from the target
or were very close and we need to be

cautious

well you dont just say thats a good
idea lets go out and do it you have to
generate a database you have to evaluate

it

retrospectively before you can jump in

so

my colleagues and i did a study of

[Music]

anticancer drugs in in
humans in our program at nci versus
the toxicity and pharmacokinetics that
were done in mouse which are primary
efficacy and sort of an early toxicity

and the histogram on the left
shows that theres a lot of variability
what have we done were just looking at

ratio

the ratio

of dose that causes toxicity

in a human to a mouse and the good news is the biggest bar in that bar chart is clustered around one that says that dose is not a terrible predictor of toxicity the dose thats tolerated in rodents may actually predict the dose that will ultimately be tolerated in humans however its far less than half of the time so we still need something else so when we looked instead again this is retrospectively because you dont do a study like this prospectively without a serious database we looked at those same set of drugs in which we looked at the area under the curve or the total exposure in humans compared to mice and what did we find we found that that curve clustered around one swallowed up some of the two tails of the curve

so area under the curve did a better job at predicting toxicity relative toxicity across species

than just looking at those ratios

the lefthand side of that histogram

um tells you the danger zone

where for some reason humans are

hypersensitive

relative to the nonhuman species that

it was tested in and so at a

at the same relative concentration
humans will have more toxicity you cant
completely eliminate that you just have
to narrow down the number of cases that

occurs

on the other hand to the right of the

tail

are the cases of where humans

can tolerate the maximum concentrations

in rodents without any trouble at all

and you can escalate

four or more times fold

that is a serious efficiency problem

its less of a safety problem

but

its the real world
so thats one one type of design its a
hypothesis that we could improve the way
first and human studies are done by

clinical pharmacology principle
that its the concentration that counts
the dose is just a way of getting the
delivering the concentration to the
individual individual preclinical

species are clinical

for those cases where there are
discrepancies and and so were sort of
satisfied with the results there but but
were curious too and we want to know

why theres a big difference
so earlier in this course sandy markey
taught a couple of courses on how to
measure drugs and how to interpret

structures and

one of the things that he didnt spend
as much time on was species differences
in drug metabolism he told you the tools
of how to develop that

but

earlier in his career he spent a
substantial time
investigating differences
in you know just take halflife as a

parameter so phenylbutazone a drug

thats no longer in the market

for humans anyway

has a threehour halflife in rabbits

six hours across a wide range of species

rat the dog

and then in humans and this is part of

the reason why its no longer in the

market for human has a threeday

halflife in humans

so the experience that you gain looking

at a rabbit a rat guinea pig or a dog

dont adequately prepare you for this

this large difference thats seen in in

humans

we know a lot pharmacogenetics tells us

a lot about the enzymes that

do the metabolism about species

differences

species variation in those we can do

better in predicting it but its always

lurking in in the background so in my

mind

metabolism is the principal compounding

factor confounding factor in first in

human studies

pachytaxal is a very successful

anticancer drug

at the time that it was being developed

all of these principles about

interspecies differences in metabolism

this would be the early 90s now were

just beginning to

sink in and have an impact on the drug

development process

and it was a little bit embarrassing

that we were about to put paclitaxol

into

patients without knowing this are there

any major

landmines in terms of its drug

metabolism

so

the folks in my lab

compared

microsomal metabolism of

in human and rat

microsomes and looked at what happened

fascinating

the

humans make a very unique metabolite which the arrow at the top

identifies this as the h for six alpha

hydroxypachotaxol

rats make none of it and rats were the

primary toxicology species for this

rats make the biggest thing they make is

pka followed by peak b and humans make

it not at all

what does that mean we dont throw out
everything but it makes you think a
little bit more carefully about how to
interpret your toxicology studies

good news is

a b and h altogether have minor biological activity for either toxicity

or activity

not so much a problem

they do influence the

amount of parent thats circulating

because the faster its metabolized the

lower its value is but it turns out that

if you just follow the parent

concentration

youll youll be in good shape because

thats where the activity and the

toxicity is

an important lesson important lesson to

learn

interaction area but heres a more
famous one in the drug interactions and
i know youve been very patient in
sitting through a bunch of oncology
examples but lets go to allergic

rhinitis

this is runny nose season right now and

most of you dont realize that the original way antihistamines work were by making you drowsy and you didnt notice

your symptoms

all right that doesnt do anything for your productivity so the pharmaceutical industry developed what they wanted to

call nonsedating

antihistamines and my former employer
the fda said why dont we just call it
relatively nonsedating and well all be
happy so thats thats what those
commercials that scroll by on the

television

the most successful the early class was uh terphenidine which was marketed as seldane was a prescription drug in the us it was over the counter
around the world and there was a pending
application at fda for it to become over

the counter

all of a sudden at the emergency room on
the other side of rockville pike in what
used to be called the naval hospital
there were a series of patients who
showed up in their late 0s early 0s
who had bizarre heart rhythms towards
side to point

and they didnt have any history of heart problems

they

had been taking antihistamines for a
while they were taking terphenidine
and it was just

really confusing to figure out what happened well what happened is they got a concomitant infection of some sort and

they started taking other medications

um

such as ketoconazole or erythromycin that interfered with the metabolism of terfenity

and it turns out that when terphenidine

0 milligrams is swallowed

what you see circulating in your body is

no terfinitine at all zero

its all the primary metabolite

fexofenidine which today of course is

marketed as

allegra or the generic form well who cares about all that stuff well

we do

because theres a onetoone correspondent relationship between

the amount the concentration of
terphenidine or effexophenidine that has
the desirable effect
and a 0 000 fold difference
in the toxicity in terms of cardiac ion

channels

remarkable

it was not important as long as every
molecule of terfenidine that was
swallowed got metabolized in the gi
tract in the liver before it circulated
to the heart

but as soon as you

then you could suddenly measure
targeting circulating in the body and
that could be related to the heart
defect problems so this was the wake up

for drug drug interactions thats a reason we have to read all these

complicated labels

and all these potential drug interactions that occur because you got to sort them out you got to prioritize

them

but

you cant overlook them particularly in patients who

take a half a dozen drugs

all right lets try one more approach

pharmacodynamic approach

so i really like pharmacokinetics helped

my career a lot but its

pharmacodynamics that what were really

interested in

does the drug work

is it good does it have too many side effects its all pharmacodynamics

and so could we design first in human trials based on pharmacodynamics instead of pharmacokinetics doesnt seem um like theres any reason not to so you have some kind of reference dose in the animals typically a maximum tolerated dose so it might be just a minimum effective dose you attach some safety factor between 0 and 0 and thats your starting dose and then you have a whole rich body of data from your preclinical studies that define the target pharmacodynamically for it if youre trying to inhibit an enzyme whats the inhibition of the

enzyme

if youre trying to tie up a receptor and block it what percent blockade do

you need

you have a very rich set of in vitro and in vivo study in the preclinical domain why not use that as a reference for assessing target impact

in humans

and could you make decisions on how large an escalation

or whether it even stopped the trial
because of futility youre already well
past your target and youre not getting
any benefit humans must be different
there are some studies like this are
done theyre more challenging but i
think the most important thing as i said
at the beginning dont get attached to
any particular design
get attached to the questions that you
want to ask

get attached to what you want to learn from doing these studies

so as as i was leaving my former
employer to come back here to nci
we were trying to internally where the
fda was a little bit out of the box

trying to

generate more innovative designs in first in human studies and trying to

enable

academic nonprofit government
investigators to do more limited studies
that would help them
take their laboratorybased studies to
the next step and so we put out a

document called exploratory inds which
essentially permitted

limited human investigation with a more
limited preclinical package
youre exposing fewer subjects youre
doing that in centers that are
incredibly talented in this area and do
intensive monitoring
essentially we were trying to
encourage molecular proof of concept
just what i was saying before
is the enzyme inhibited is the receptor

blocked

and we were also trying to encourage functional imaging which had been sort of caught in in a difficult space in the regulatory world and needed to be well defined in terms of that can be a generator of biomarkers and can be helpful as an exploratory sense as well

at nci

my soontobe colleagues were working on something called phase zero which was very much the same thing as lets up front limit how far were going to go into the clinic in return for

getting there faster

so as as we saw earlier in this talk the historical phases of human evaluation phase one is about safety safety safety and itd be nice if you could learn something about activity you should certainly look

youre likely to find some surprises

phase two is where youre youre really

making your business model

is the activity promising or is it not

can i invest in this compound

or is a five percent response rate too

low

or is it too toxic

and then phase three is comparative is
does this compound does this drug
development program have anything to
offer thats an improvement over current

therapy

phase zero is stepping back from that it

doesnt exist as a

real entity to exist as a concept there are phase zero studies done ill show you one of them but its really about what is the most important thing about

this compound that you want to learn as

soon as possible

everybody would say does it work

well yeah but you got to back off from

that youve spent

millions of dollars figuring out why it

works selecting the thing that has

the best impact on

a particular

mechanistic target

you want to know

whether you actually get the desired

effect you dont get that by just

measuring whatever the dose limiting

toxicity is

or counting the number of responses and

dividing the number of patients that are

on those are incredibly important

endpoints in the trial

but they come later

and to get higher doses require

more safety study preclinically but you

could learn a lot about mechanism of

action you could answer important

questions

if you had a process like an exploratory

ind or a phase zero

i cant tell you

how hard it is to get people to articulate the question they want the

answer to

i would have thought

everybody can knows what they want

they want to go to the next stage

is essentially what you know i usually

hear what is the key piece of

information some of the sometimes its a

very limited question

that builds on experience with analogs

that had

problem getting in the body or a problem

with toxicity or something like that

but when you really have a novel

chemotype

what is the what it what do you really

want to

do and how are you going to figure that

out

so almost all drugs are given orally

and

they dont work systemically unless they

get absorbed

[Music]

they dont always get absorbed
a fair number of drugs are chemically
unstable at ph

in your gi tract
or even unstable if you take proton pump
inhibitors and have ph
theres also a lot of enzymes that line
your gastrointestinal tract
and every molecule essentially
has to go through the liver before it
goes to the systemic circulation
and then theres those formulation
things you swallow a nice
hard pill and it comes out the other end
as a nice hard pill that actually does

happen so

a key question is

this is my drug this is the thing that i

think has the best chance of working

can i achieve adequate concentration so

thats simple thats not very

complicated

why not do that in your first set of

patients

in a project we did in this building
in on the nci clinical service we were
looking at an enzyme
inhibitor and we wanted to know whether
this dna repair enzyme was inhibited
we couldnt tell that from looking at

look at it

bioavailability studies by looking at

whether it was absorbed we could only

by measuring the products of the enzyme

reaction

all right

did we do it or not

were not interested in whats circulating in the plasma this is pretty

tough

were interested in what happens
selectively in the tumor and so we have
to look in the tumor pharmacokinetics is
hard enough when you collect
plasma and urine and stool and
occasional opportunistic samples
somewhere else but to design a study
that requires a tumor biopsy it requires

really

patient population

and dont do it without an incredibly

welldocumented assay

you know you dont develop your assay
in tumor biopsies you have your abs

assay ready before you get there

so what happened upstairs on the th

so valiparib was a parp inhibitor that
we partnered with abbott pharmaceuticals
on we did the first in human study here
it was a phase zero it was only intended

floor

to be a single dose
a single dose is not going to be
therapeutically effective because you
need to inhibit the dna repair for some

period of time

its unlikely to be toxic as well so the

fda was fine with us choosing this

mechanism of action to go

so we started at single doses of 0

milligrams went up to 0 milligrams

and by 0 milligrams we had reached the

target what was the target

other than wanting to make sure that it

got in the body at all

we looked at the concentrations that

were circulating

in the preclinical studies

that

of tumorbearing animals in which there

was a target effect

so essentially we were saying as youve

learned in many lectures in this course

that the concentration circulating in

plasma can give you some information

about what target youre looking for

this is not the final answer but if you

cant do that then

the rest of the rest of the

process is going to be more challenging

so we met the first goal what about the

second goal and thats

can we get definitive results out of the

tumor tumor sample

and so heres a series of a half dozen

patients that were were studied again

here in this

in this building um in the first in the

phase zero study

and for five out of six

at that dose that achieved adequate concentrations based on preclinical studies for five out of six

pretty

much depleted entirely the product of

the enzyme

so we had a dose that
looked good based on preclinical studies
and based on actual studies in tumors

we had one

looks like it intended its goal

tumor that barely responded
that just shows we didnt make up the
data because thats the way real real

data actually works

are we finished

should we invest a billion dollars in this drug not now theres still a lot to

be done

it could be that theres a disconnect
between the mechanism of action and the
actual clinical antitumor activity a
lot of work to go but youre willing to
make larger investments when you have
these kinds of data

we did a variety of studies by looking
at different doses and different time
intervals between
the dosing to help us define
how to do what essentially was
almost going directly to phase two
because we were we were ready weve had
proof of concept we were ready to see uh
see what could be done
we needed to know a little bit about
those ranging and um

dose intervals

let me say something about uh functional imaging which was the other part of this guidance um bobbiness gave a talk he was like the third one in the in the series

uh in the fall

and he covered this from the standpoint of pet as a tool more to learn something in particular about the brain

and

um

very very attractive

tool were greatly embedded to the
neuroscientists for here at nih and
elsewhere for developing it but in terms

of new agent development
lets think about the questions again
i want to know if the treatment impacts
the desired target if it doesnt block
the receptor im not going anywhere else
okay so

somebody appointed me the manager of a

billion dollar company to

figure out what things to go im not

going to go further if it doesnt impact

the target its too early

and everything i invest after thats

going to be very high

i do want to know what the minimum dose

it causes in effect
and i want to know what the maximum dose
beyond which you no longer get any

effect

the shape of the dose response curve
and my marketing department has said
that this pill can only be given once a
day because thats what patients like
but i need to know scientifically
whether once a day is the right dose
interval or not and
its very clumsy to do this in just

empiric studies of toxicity and activity
can be done has been done is done
but maybe if we had biomarker driven

[Music]

answers to these questions and maybe if we use the pet technology we could find

some ways of doing this

with just a handful of subjects

so this is a different version of the

slide that bob inna showed in his talk

back in the fall

which im going to use to address these

three questions so the first question is

so that i can tell that nasty manager

whos analyzing

managing the pipeline

this is an enzyme

monoamine oxidase type b

its a

reversible inhibitor losabamide

investigational

did it do anything to monoamine oxidase

type b and you dont have to go to

medical school or nuclear medicine

school to say

that at milligrams

twice a day in the upper right hand
corner there is far more bright spots
than there were at the baseline scan in
the upper left corner

furthermore so yes the answer the first

question is yes

the second question the most minimum maximum doses are whats the shape of the dose response curve and thats the lower lefthand current when i double

the dose

the few remaining flickers of candles that are left in the brain

disappear

ive got complete inhibition im wasting my time going any higher assuming this

is the mechanism

the third one is the challenge is can i give this drug once a day or is it going to take more often it was given twice a

day in the um

[Music]

in the investigational study because the the established marketed agent depornell

is

is given twice a day so they thought

they would use that schedule

the pet image was taken hours later

what does that say that picture is

not quite as bright as in the upper left

but its pretty much there

it looks like hours would in addition

to being impractical would be terrible

it doesnt tell me whether i need to get

i know that every hours works because

we gave it the idea and we looked at the

end okay

so somewhere between works and

hours doesnt

is the answer to whether i can give it

once a day

these day this this design

this experiment didnt answer that

question but if we had had a hour

experiment it would have

um

fascinating approach only takes a
handful of subjects to figure out
whether whether youve achieved your
molecular goal or not does this say
whether loseabamide is going to be a
blockbuster drug or not

according to published results
lizebramide was pulled from clinical
development because of hepatotoxicity
all the brain scans in the world folks
arent going to tell you anything about
hepatotoxicity and
you can minimize your risk but
youre always looking
so to sort of get up to the wrapup
portion

[Music]

when i first started giving these talks
many people were concerned that first
and human trials were changing right
before their eyes that there were
different questions that were being

asked

everybody was under great pressure to do them faster

there are all these laboratorybased correlative studies whereas previously phase one studies have been very simple well now first in human trials arent necessarily phase one theyre also phase zero

and i think that the community who does these studies is now comfortable with

that so yes

theres an identity crisis if you think
that your design will work the same for
every drug that comes off your assembly

line

but

the most important thing is youve been challenged to figure out what design you really need for your drugs so maybe its an identity crisis but its good the other thing and really the last take home message is that that is invariant regardless of what design you use is no matter what design you use whats going to be inherent in first in human

before

studies if its never been in humans

theres a good probability that youre going to see something different in

humans

that you werent prepared for in all the nonclinical studies that you

sometimes its actually a pleasant

surprise

sometimes its annoyance its a scale

you know one end is pleasant in the

middle is an annoyance

and on the far side is its a deal

breaker

a side effect that wasnt predicted

by animals a

molecular biotransformation that wasnt

anticipated

a parallel pathways that overcome

the intended mechanism of action

this is an excuse not to do the best

design you can

but its a cautionary tale that you

always keep looking

and clinical pharmacologists by nature

are often the first time that first

people who regardless of the discipline

who ask questions like well what

happened to the hearing

what happened to the hearing yeah

theres some related molecules to do

that or

hows the kidney function all those data

are there but my goodness theyre piles of data

you have to look at it you have to make sure you find the surprises when they happen uh lastly i just thank uh my colleagues uh here at nih and and my former colleagues at fda for um helping me learn a lot of this stuff participate in clinical trials and the yellow folks around the edges of this diagram reflects some of the sites of people whove signed up and taken for this lecture they also are representative of the people that ive collaborated with outside of the nih at the f and the fda and i thank all of them for helping me pursue this interest