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director of the departolo family
personalized medicine institute at
moffitt cancer center
dr mcleod has over 0 years of
experience in pharmacogenetics
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pharmacology

his research interest focuses on
pharmacogenetics and the role of genetic
differences in individual responses to

cancer drugs

after earning a doctorate of pharmacy
degree from philadelphia college of
pharmacy and science dr mcleod went on
to complete clinical research fellowship

at st jude

and a research fellowship at the
university of glasgow

please enjoy dr mcleods lecture today
hi im dr howard mcleod im going to
talk to you today about dose

modifications based on pharmacogenetics
research

this is work taking the human genome and
trying to make it so that more rational
therapeutic decisions are are being made
over the course of this time well talk
about examples

that are influenced influencing dosage
selection of therapy and other ways that
the genome can be useful in this manner
now lets just start off with a quote
that a surgeon who uses the wrong side
of the scalpel will cut his own fingers
and not the patient

if the same applied to drugs they would have been investigated very carefully a long time ago this quote is supposedly from 9 but its very relevant today of the medicines that we have available we know something about them enough to help many people but theres much work still to be done to understand why

theres variation

across populations why does one person
have a benefit from a medicine someone
else have a side effect from that exact
same medicine someone needs a big dose
someone might need a small dosage of

that same medicine and so well be exploring that a little bit more in the in the following section now the clinical problem is a really wonderful problem and that is that for most therapeutic areas there are multiple different active therapies that are available to treat that disease if there is only one medicine there wouldnt be a lot of choice we would just hope for the best but there there are choices and so how does one select from amongst the available medicines in order to treat a patient we need to be taking into account the likelihood of benefit to response to therapy the the chance that theres a bad effect an unpredictable toxicity that could

occur

and then theres also an issue that many
of us like to ignore
and that is that medicines cost a lot of
money many of the new anticancer drugs
can cost as much as twenty thousand
dollars a month a wellinsured patient

still has a major financial burden uh
even though they thought they were
covered by insurance and so we need to
be able to make rational choices in
terms of benefit
of toxicity and also making sure the
patient can afford to get the kind of
treatment

that is

needed for their disease whether its through a societys financial burden or the individual patients financial burden and so we need better tools to make those choices as we go forward now there are many different factors that can influence a medicine shown on the outside of this wheel are a number of the different factors outward factors that can be important there can be things like stress liver function kidney function sunlight many different aspects of of our daily life can be important in terms of influencing the way drugs are metabolized the way theyre absorbed other features of a medicine

at the hub of the wheel though is our genetic constitution and so the way that our dna is coded influences how we look

influences how we look

how we taste things how we smell things

often even the way we walk can be

influenced in part by our dna

that also could influence things like

the metabolism of a medicine or or how

long it will stay in the body and so we

look at genetics as one part of

understanding how to pick the right

medicine and the right dose of the

medicine for an individual patient

now when we look at the human body there

are two different factors that we like

to talk about

involved in pharmacology of a medicine

in a person

the first is the pharmacokinetics and you can think of that as what the body does to the drug so a drug is absorbed

it is metabolized by the liver its
eliminated by the kidneys those are
pharmacokinetic factors and so dna
changes in those factors can be

important and well show some examples of that

also theres the pharmacodynamics you can think about that as what the drug does to the body so if a drug binds to a certain receptor in this case on a on

the

surface of the lung uh an an inhaled
albuterol or cell butymal and as its
known in many countries will bind to
this receptor cause the airway to open
and relieve some asthma uh symptoms so
this idea that a drug hits a receptor
and causes an effect is also an
important feature and so there are
examples where genetic variation in both
pharmacokinetics and pharmacodynamics

can be important for
influencing how a particular patient
will respond to a given medicine
now when we look at the genetic factors
that influence drugs or pharmacogenomics
pharmacogenetics many different terms
that can be used there are many
different examples that are approved by

the us fda

now there are over 0 medicines that have genetic information somewhere in the prescribing recommendations that are put out by the fda but this list thats shown here is a subset of those where in the dosing and administration section genetics as a is a feature now some of these are genetic abnormalities that are seen in a disease like cancer so gene amplification gene deletion some other gene abnormalities will be in the in the tumor not necessarily in the normal tissues but then many of these uh of these examples uh that are shown here are shown in the normal dna and they can influence the dose of a medicine the toxicity risk for a medicine many different features that one can take and

use

in terms of guiding
the therapies that one receives so
pharmacogenomics is not something that
might happen someday
but rather there are many different
examples of commonly used medicines
that that for where this is important

and this includes some of the medicines
that we use across many different
diseases

controlling pain uh preventing vomiting
using antidepressants uh adhd drugs anti
blood thinner or blood thinning drugs or
anticoagulants

um its not just for cancer and and the
markers that might be in a tumor but
rather features that are there so
someone getting a surgical procedure no
matter what its its for will have pain
issues if theyre getting general
anesthesia will need to have an
antivomiting medicine
someone that is getting hot flashes from
an antiestrogen or going to just the

the

for the hot flashes

or might need an antidepressant for the
treatment of depression or they can be
used for chronic pain many different
ways that these medicines can be
important and so its not just the
highly specialized diseases

where genetic variability has become important but really very commonly used medicines

are being influenced by by genetics now there are guidelines that have been put out to to help us use these medicines so theres a consortium called the clinical pharmacogenomics implementation consortium cpic as its known that has been put together its based in the united states but has representatives from uh over 0 different countries there are about 00 different institutions involved and so we we get input from many different bodies on how to use these medicines in a in a safe and effective way if a patient comes in with a genetic result how can we use that to guide which therapy which dose of therapy in some cases even what route of therapy intravenous versus oral that one might use and so these guidelines are very useful and they go across many different drugs and many different disease areas and im not going to list them all for

you you can go to the pharmacogenetics
knowledge base and and see the list for
yourself

but

showing that this kind of information is available and can be useful for guiding your health system guiding your individual practice
guiding your university whatever the setting is that youre thinking about this these guidelines can be quite

also theres an opportunity to be involved in this if youre in a country that does not have these guidelines and you want to be involved with cpic

useful

guidelines

um reach out

for for that because its a great
opportunity to get involved and help the
usage of of these medicines
now there are many ways that precision

pharmacogenomics pharmacogenetics is a part of that can be used for for thinking about medicines one is to avoid

medicine or

a medicine that might have a very bad side effect and the presentation from dr pakinowski a few sessions ago highlighted some of the allergic reactions that could occur where you would want to avoid that medicine at all costs because of the severe nature of

that abnormality

you to know that a medicine is more
likely to work and therefore should be
included in the options

explaining a bad side effect so if a
cancer patient has a drop in their white
blood count after a chemotherapy drug
you want to know is it that medicine or
is it one of the other medicines the
patients taking thats causing that
drop because you may need to use that
medicine but if theres a severe
reaction you may need to avoid it
because so doing genetic testing to

explain

is it a genetic abnormality for this

medicine or for another one it is

certainly a part that can be can be can

be done but im going to focus in on some examples

for choosing the dose of a medicine or choosing from amongst a number of available medicines where you need to pick the medicine and then the dose in in the uh over the next little period

of time

now one of the examples that was shown by dr pakinowski during his presentation

was a medicine for
attention deficit hyperactivity disorder
called atomoxetine this is an example of
a drug where in the patient prescribing
information their socalled package

insert

a gene called sipd is listed and

therefore

and anything in the metabolic pathway
thats influenced by sip d
can be influenced by those who have
genetic abnormalities so a small
percentage of the population have extra
copies of this gene
about 0 percent of the general
population in the united states have uh

are are missing this gene and then
theres others that are in the middle
and so you have some people where they
would need different doses based on
their genetics and so within the fda
prescribing recommendations it mentions

the

sip d poor metabolizer status how that would increase the result in higher blood levels higher rates of adverse reactions and recommends different dosing based on both body weight and genetic information so heres an example where environment in this case the patients body size and genetics in this case their d status both will influence the dose thats needed and the the frequency in which one uses the medicine now look at the changes here a half a milligram per kilogram per day

increase every three days

if they have small body weight

if they have a larger body weight 0

milligrams per day not including kilograms so so a very different way of dosing and then if theres a genetic deficiency instead of every three days its every four weeks that one would make these changes very dramatic difference in terms of how one would manage this medicine for a child with this disorder and as you can imagine children with this disorders have a terrible time in terms of their education in terms of their social interactions giving this in the wrong manner causing them additional side effects or not giving them the efficacy they need could really have a terrible effect on this child and the in the way that

they can uh

be part of society so its an important factor and an example where the fda has made those those important changes now one of the older examples thats quite important is with the socalled thiopurine medicines these are medicines as a thioprin mercaptopurine guanine or

the ones that are available in most
countries that are used in the context
of a solid organ transplantation so
heart kidney liver transplantation
also in the treatment of leukemia in
particular childhood leukemia and these
medicines azathiprine is a prodrug
which is metabolized

to mercaptopurine

which then can either go and form uh an active metabolite thioguani nucleotides

as theyre known

or can be inactivated by thiopuri and methyltransferase or tpmt is as easier

to say

and so theres a competition between
whether this medicine is activated
or inactivated based on the the genes

that are involved

and so as you can imagine genetic

abnormalities that influence the

function of this gene will have an

importance on how much medicine can be

tolerated for a given patient and so if

you look at this slide heres a general

population within a population there are

some people who have the the normal two copies of the gene have normal functioning activity there are some people who have one normal copy and one abnormal copy copy and so i have an intermediate ability to break down the medicine and then there are the the less common individuals that that have uh are either a low activity or are completely missing the gene and therefore have very little ability to break down this medicine now if you give all of them the exact same dose what youll find is that some people have standard levels of in the of the medicine in their blood those that are missing one of the copies of the gene will have a little bit higher levels

those that are missing both copies and have no ability to break it down or an effective ability to break it down have

much higher levels

and then what that translates into is a

normal risk of myelosuppression a drop

in white blood cell count

an intermediate risk or a very acute and high risk 00 percent risk of getting severe toxicity many of these children need hospitalizations um there if its an organ transplant it may result in in other bad effects terms of losing the

kidney or the liver

and so very different scenarios between

acute toxicity or very manageable late

occurring toxicities that can occur in

instead

these patients and so if one goes and

uses the genetic information to give either a standard dose

a reduced dose

or a very reduced dose based on normal intermediate or absent enzyme activity one will see that the blood levels that are obtained are fairly even across the population because just a little bit of the medicine in a low or absent metabolizer will result in quite a substantial blood level

that also means that everyone is at a very similar risk of a drop in their white blood cell count and at least in

the context of leukemia

the survival studies have shown that
using this method individualizing the
dose based on genetics resulting in even
an even amount of blood level
even amount of toxicity also results in
a high level of cure of these children
with leukemia so whereas the patients
who are getting are absent for the
enzyme and giving a standard dose have
severe toxicity cannot get their full
therapy and have a higher risk of

relapse

in this case high chance of cure
moderate chance of toxicity a much
better scenario for these very severe
significant diseases that are being

managed

now you can look at cpic guidelines and
see that there are specific
recommendations for each of these
medicines depending on whether you have

normal activity

intermediate activity or low or absent activity where you would make a a substantial reduction down to 90 percent

of the normal dose or
only less than 0 percent i should say
reduction by 90

just given three times a week compared to a full dose given every day in those with normal activity so a substantial difference in the use of these medicines in order to achieve the same high level of success that is that is influenced by the the genetics now these children cannot be ascertained by their outward appearance the way they walk their favorite foods theres nothing about them that would tell you that they are missing this gene or have normal copies of this gene its only at least in the old times when you gave the medicine and something bad happened that one could figure this out now genetic testing can occur prior to administering the medicine the dose of the medicine determined based on those results and those good outcomes are more likely to happen because of this sort of approach so an example of how this

this genebased approach

pharmacogeneticsbased approach can be used in the context of therapy now a similar finding

for

another cancer drug called tamoxifen
this is a drug thats been used for many
years in the treatment of estrogen
receptor positive breast cancer its an
important drug for curing breast cancer

its been responsible for saving many lives over the decades but it has complex metabolism where this drug is metabolized to this active more active metabolite called indoxapin and this enzyme sipd that we already talked about earlier is an important factor there and so as i mentioned before 0 of the population are missing this gene cannot functionally activate tamoxifen at the same rate that the rest of the population could and so one could go and and decide what to do based on that now there are studies showing a relationship between sip d genetics and recurrence of breast cancer there are also some studies where that

association was not observed so its not a clean story where every time the genetics will be important but in most of the studies using using standard dose tamoxifen a difference has been seen where a patient that has two normal copies a socalled extensive metabolizer will have a much better outcome than a patient with only one normal copy or the patients with no normal copies of the gene so you can think of this as a lot of activation of the metabolite active metabolite a moderate intermediate activation of the metabolite or relatively low activation and theres an impact on survival in the treatment of breast cancer based on these results

so

what can one do for that so a clinical trial was performed its been was published initially back in 0 as ill show you in the next slide theres some data that has been published in late

0

but this looked at a large number of

patients initially 9 patients
subsequently published over 00 patients
where they took the patients if they
were extensive metabolizers so two
normal copies of the gene they gave them
the standard 0 milligrams per day of

tamoxifen

if they were genotyped and had an intermediate metabolizer status so one normal copy one abnormal copy they were given a doubling of the dose 0

milligrams

every day now the fda approved dosing is

between 0 and 0 milligrams so this is

all within the fda approved dosing

but a doubling of the dose and what you

see is that starting off after four

months of the 0 milligram dose you see

a substantial difference in the amount

of active drug available in the

when you then keep the standard dose for
the these patients here and give a
doubling of dose for the intermediate
patients you now cannot tell the
difference statistically between these

patients body

two groups youve had a normalization of
the blood levels based on a
doubling of the dose for some patients
and a standard dose for the rest of the
patients so this idea that we can use
genetic information

from the start of therapy

choose the right dose of a medicine
and then administer it is really

important now in the example i just gave
you with thiopuri methyltransferase and
the thiopurine drugs

you had an intermediate biomarker you had toxicity as a factor that could be

measured

in the case of tamoxifen there is no intermediate biomarker you cant tell the drug is working or not working unless a patient has recurrence of their

disease

and so this is a high stakes endeavor
you want to give the right dose from the
very start in this context and not give
a dose that you hope is the right one
and then find out later the patience has
has recurrence of their disease and so

its a serious matter

and not one where theres another way of managing this and so we need to have

this sort of information

now i showed you a complex metabolism

of tamoxifen in a previous slide

and so one could then go and look at

other factors that are important in

terms of metabolism in terms of other

outward appearances and a study just

came out in october of 0

where a group went and looked at 00

women that got tamoxifen for the

treatment of breast cancer

looked at sip d but also looked at

other genetic factors such as acepc9

sorry sepc9

and then about 0 other genes were also

evaluated looked at body weight they

even looked at what season a patient was

treated and what they found is that the

other genetic factors and the other

clinical factors

were not significant in terms of

statistical values for the

the dose of of tamoxifen needed to get

the adequate blood levels but sip to c9 metabolism

had some effect

body weight had some effect for reasons

that are not

yet clear

season also had some effect so were trying to understand why that is and see whether that can be used in terms of of uh treatment also can be used in terms of explaining the mechanism but the biggest factor was sip d and so using sip d supplementing it with body weight with some other genetic factors certainly can be a way of trying to normalize this treatment and theres just an example of trying to not let one factor be the explanation our bodies are very complex the metabolism metabolism of medicines are complex its no surprise that its going to be multiple factors that come together and influence the the uh way a drug is handled to get the achieved goal and so we cant

think of life as being too simple we

need to be

embracing complexity and trying to
understand it um and that will allow us
to get to the point where we can really
dial in the right drug the right dose
for a given patient

now

one last dosage example i want to give
is for an antifungal therapy in this
case in the context of leukemia where
where fungal infections can be fatal
and a gene we havent talked about so
much yet called cypc9
now as many of you know these genes
are cytochrome p0 genes therefore the

sip

and then they were numbered based on when they were discovered and whether they were part of a family member so the second family to be discovered the the uh third branch of it or socalled c branch and then the 9th member of that family um is why its sip c9 now its not very easy to remember necessarily but there are only a small number of genes that are are critical

for the use of pharmacogenetics in patients and so it is one of those that is worth remembering now as i mentioned invasive fungal infections

are a major contributor of morbidity
prolonged hospitalization and mortality
in cancer patients who have dropped
their white blood cell count and
leukemia is a prime area where their
white blood cells are wiped out based on
the therapy so that normal cells can be

reconstituting there and so we we give

antifungal therapies as prophylaxis you
give the therapy to prevent the
occurrence of a fungal infection not you
dont wait until someone has a fungal
infection when youre then trying to
chase the infection and hope that the
patient doesnt die first and so
voraconozo is one of the the many
medicines or the one of several
medicines i should say
that are effective

preventing fungal infection or at least
minimizing the chance of fungal
infection in the context of of acute
leukemia

and this gene cypc9
is one of the genetic variants that can
influence the blood levels of of this
medicine and so if you look at this
medicine its metabolized by several
different genes but sip c9 is the the
predominant one that can cause a
inactive uh version of that and there
are are variations in these genes such

that can influence that now other
presentations have gone into some of the
the nomenclature for these genes but
with each variation that has occurred a
new number has been assigned to it its
basically a way of trying to keep track
of these variations uh its its uh easy
for those of us who work in the field to
remember sip c9 star

as the star variation

field

but certainly if youre entering this

it is complex figuring out what does

but

these genetic variants as you can get used to them

will have meaning in terms of their application now when you look at these genetic variations and look at blood levels as i mentioned those who have

a are

extensive metabolizers are shown here in the middle those are the ones with normal amounts of the enzyme activity those that are are poor metabolizers theyre missing both copies of the gene have very high blood levels and so you worry about toxicity in these patients being caused by the medicine and then those who are ultra rapid metabolizer have extra copies of the gene they have lower blood levels and in many cases can never get to a therapeutic blood level of these medicines and so one can look at a number of variations i mentioned the poor metabolizers that would be in this category here the people who have the

socalled star either together or in combination with star now star star can occur but is extremely rare in most world populations but theyre at risk for for extra high blood levels and the toxicities that could occur there but the ones that were concerned about in terms of preventing a fungus infection in in patients with leukemia are either one copy of this variation or two copies of this variation because the star variation causes extra amounts of the protein to be produced and so the body gets rid of the medicine in a much more rapid fashion and so

these folks

have a very low chance of getting the right blood levels their risk for low blood levels or extremely low blood levels and thats a major concern in terms of preventing a fungal infection

and so

these are the patients where its now routine many centers to genetically analyze these patients prior to starting their therapy their their antileukemia

therapy so that one can get them on the
right antifungus prophylaxis from the
very first dose
rather than trying it out and hoping for

also

the best

an important feature that i wanted to include here is that when you do genetic analysis this is from a paper uh from 0 that looked at the cost of screening all patients for this genetic factor looked at the blood adding blood levels to to help manage these patients looked at how many uh fungal infections would be avoided and with uh the lowest amount of savings still were saving

about

dollars per patient by doing

preemptive testing so in a group of 00

patients a health system would be saving

in this case and

over 0 000

avoiding a severe lifethreatening or fatal toxicity in the patient and so not only is there a strong clinical case for doing this sort of testing but also a

positive financial case for doing this

and so two fantastic reasons to be doing

a genetic test to try to optimize the

therapy

in this very lifethreatening situation
theres also an as an example
dosing algorithms or workflow diagrams
that can be conducted so in this case
you want to use voracons or prophylaxis

if theyve had

a bone marrow or liver transplant then
using white blood cells to do the
genetics might be measuring someone
elses genotype either the bone marrow
donor or in this case the white blood
cells would be from the patient the
liver and the liver transplant would be
from someone else and so you need to
take a different route there but if its
a leukemia patient can be genotyped
based on the results either the standard

a uh higher dose of oroconazole or a switch to a completely different therapy um can be can be the result

dose of voraconazole

based on that and so a difference in dosing a difference in therapy based on the type of results one gets and we can know this prior to ever treating the patient because of rapid turnaround genetic testing and so is an example again where pharmacogenetics influences the dosage or the choice of therapy and by doing it preemptively you can prevent that sort of uh effect from occurring and not have to play catchup when a patient has a terrible infection and is uh is at risk for dying from that

infection

now i focused in on examples where pharmacogenetics could influence the dose that one might pick for a patient the last example hinted at a little bit of selection of therapy and so want to spend a little bit of time on selecting from a menu of available therapies

for

one really important area for society

for this sort of work is in the

treatment of depression and theres two

major classes of antidepressants both of which have some pharmacogenetic data on which to influence it now the first class is the tricyclic antidepressants these are the older classes of medicines but but again very very effective um they are our names are shown here um and they can have pronounced uh serotonin or neuro or norepinephrine or adrenergic effects and so dry mouth some of the other side effects like that can occur with these medicines that wouldnt occur with some of the newer medicines but these medicines are used for depression also can be used for chronic pain and such as trigeminal neuralgia and other syndromes are like that and so its important that we understand can genetics influence the the dose of these medicines or the selection of these medicines and so one can look at metabolism and amitriptyline is metabolized by this enzyme to nortriptyline its inactivated by this other enzyme to this metabolite here so two different genes you can see

are already important in terms of what sort of side effects and what sort of efficacy one might get for from these classes of medicines and and certainly one can expand out by going to this example this is the 00 0 paper theres a new a new version that will be coming out in the future of the cpic guideline looking at these two in gene these two genes and the doses of tricyclic antidepressants that one could use for the treatment of depression or other disorders and so for each of them there is metabolism by sip d and the dose reductions that would that could occur metabolism by sip c9 the dose reductions or avoidance that one could occur and and so thats important now i want to also mention that genetics is not the only way of trying to individualize

medicines

there have been

examples over the years where these medicines can be have their blood levels measured so a blood level taken a blood

sample taken plasma or serum
removed put onto a machine that can look
at how much medicine is in the blood per
unit and and one can look and see a
toxic range or a therapeutic range for
these these medicines and certainly if
you have the access to these these types
of blood level measurements in a timely
manner its a very viable and useful way
forward in terms of choosing the dose of

medicine

however many of these assays are not widely available or are not available in a timely manner and so waiting a month for a result is usually not the ideal for a given patient you dont want to just try one of these and see what happens you want to be able to

understand

whats going on also with the genetic information especially with a rapid turnaround assay one can understand which of these medicines are a very low yield a very have a very low chance of benefiting the patient and therefore can be avoided altogether and which ones at

least have a chance of benefiting the patient and can be tried out and so having the information prior to ever giving the medicine gives the best chance of of having that sort of thing now if you start the patient on the medicine you can then get blood levels and see if theres some finetuning that can be done but i wanted to mention this because blood levels therapeutic drug monitoring is a viable way of trying to individualize therapy it just cannot be done prior to administering the dose but rather the patient has to be on the drug for a given amount of time

now a second class of these medicines
are the selective serotonin reuptake
inhibitors a little bit newer class of

medicines

they have a similar side effect profile

but a little bit less frequent

occurrence as the tricyclic

antidepressants

and uh therefore

and response rate has been associated with genetic polymorphisms in these same

genes that i mentioned before and well
get into some of the response data
in a in a few slides uh but uh its the
same genes that are important and again
cp guidelines have been produced
have been important in terms of putting
to pulling together data on how should
these medicines be used and in what way
these medicines are metabolized by
multiple genes in this case with
peroxidant

its subd hip is also featured in

the others

sypc9 is important here

whereas c9 is not important for these
other medicines and so you get into some
some ways of of trying to look at these
these drugs also when you look at the
cpic guidelines and the the online
supplements that are available for them
you also see a lot of the background
data in this case is for peroxison and
septuc sipd you can see
many of the clinical trials and theres

also a rating on the quality of the data

this trial here

or these set of trials had a very high
level of evidence uh this study here was
a relatively weak uh study might have
been because it was a case report or
small number of study a small number of
patients in the study or some other

factor

like that but this sort of consensus
around the studies their level of
evidence how they can be used is an
important resource thats freely
available for anyone that has access

to the internet

and so these guidelines again sticking
with proxetine as an example the
guidelines will make

recommendations on if you have a ultra rapid metabolizer so you have an extra copy of the sip d gene what the

implications are

whether

a different dose should be given or in this case ill really select a different

drug

that is not using this enzyme and the

data for that is quite strong

and so

this type of information is quite
helpful in terms of developing local
guidelines for practical application of

this work

enough you want to know the dose or
whether a drug should be avoided and
thats really a key component of these
these guidelines and so in the case of
paroxetine it has a specific percentage
dosage reduction for the poor
metabolizers or avoidance of the drug
for for this other drug fluvox of

fuvoxamine

a reduction in dosage
or avoidance no dosing recommendation
for fluoxetine those studies have not
been performed therefore we couldnt
make a clear recommendation and so this
sort of information is there

consensus of multiple different investigators from multiple continents to try to pull together recommendations

publicly available

that you could use
as youre trying to use pharmacogenetics
to dose dose patients
same with these other medicines in these

same categories

so that sort of information is important but when you look at the utility theres also other studies that have been performed some of these studies like this first study looked at whether testing and acting on the testing would reduce cost and in this case it reduced cost quite substantially probably mainly because of the longer the the differences in hospitals stay but its important in terms of the different costs that can be there in some cases testing reduced the cost by a blanket amount in this case testing was done but not no intervention happened and those who would have been intervened on had a much higher health

care costs based on so you can pull

together individual examples of
literature that really start telling you
theres differences in outcome
differences in cost and its important
so for example this study that thats
shown here um looked at patients who had

been genotype

genotyped versus patients who had not been genotyped this was not a randomized study but rather looking at at patients at a individual center they looked at sip d and sypc9 and what they found is a reduction in overall drug costs during the first year of this evaluation and it was almost a thousand dollars a

difference

in in terms of the the the costs that
were there um enough to more than pay
for the testing uh for for these these

patients

additional

more more

carefully guided studies have been done in this case looking at standard use of

of

antidepressant therapy versus genome

guided use in the red

of these therapies and they looked at a

number of the different ways of

measuring success in terms of depression

these are different uh different tools

uh surveys other questionnaires other

that can be used to gut to see is there
a benefit happening and then one can
look at those those effects what you see
as no matter which way you measure the

clinical tests

the

antidepressant benefit it was higher in
the genetically guided treatment than it
was in those who got the standard
clinical management alone
with using other genes not just
metabolism in this case a target one of
the receptors im looking at gene
polymorphisms you you see a a much
better plateau of the of the uh
depression results in uh and much more
quickly uh in the patients um who had a
particular genotype compared to the rest
of the patient just i identifying or
highlighting that response

can be quite different

in these patients

occurring much more quickly and in a

larger number of patients based on the

genetic factors

now this is important because you dont

want to have to go out weeks

to then be able to tell a patient that

the medicine didnt work for them you

want to know early and so by optimizing

the the chance of benefit using whatever

genetic tools are are

most relevant and most most effective

is an important factor in terms of how

one uh really starts treating these

these patients

now there was a recent study that was a

randomized doubleblind clinical trial

um some of the features are shown here

in terms of of what was done in terms of

interventions um you can see some of the

features of the patients theres about

00 plus patients that were randomized

fairly evenly to the the genetics group

or the clinical management group show

you some of the factors theres a whole

table of other factors where there was
no significant difference between the
two treatment groups and what you see
here

is the response rate

it was measured by a number of different
factors i chose one of them from figure

of that paper shown here

was was much higher statistically
significantly higher and clinically
significantly higher in those patients
who got the gene guided therapy versus
those who got standard clinical
management

now an important factor is that the the severe toxicity
interventions the burden of toxicity as

its called

was was

much

better controlled in the genetic guided
therapy compared to those who didnt get
genetic therapy and so its one of those
situations where better efficacy
lower chance of toxicity was occurring
with genetic guided therapy compared to

what you see with standard clinical management

now most depression is managed in a busy

primary care environment its its not

that common that a psychiatrist or

psychologist is involved in that

management and so you can manage how you

can imagine how in a busy clinical

environment

understanding the genetic risk of

therapy

choosing the therapy based on the best chance of benefit lowest chance of

toxicity is going to be

a much more efficient way of managing patients both in terms of clinician time and in terms of the patients time a need for return visits etc and so were seeing a lot of use of these types of testing in the context of of

of depression

trying to improve that and theres been data now for other types of psychiatric

care

that that can be managed and this slide

here is just trying to remind that

theres going to be a genetic
information pharmacodynamics
pharmacokinetics but then other factors
age the severity of disease the
environment both the natural environment
the home environment the work

environment

other factors depression schizophrenia other mental health disorders are very complex disease its not just a simple gene defect and so genetics will help

but will never be the only tool
for use in in guiding therapy and so
this is just a slide to remind myself

guide the therapy

and you

that genetics is a a factor that can be
added to good clinical management
not a replacement for good clinical
management a robot is not going to be
able to take genetic results and manage
depression it still takes good clinical
management but hopefully better results
for the patients better efficiencies for
the practice as we go forward

weve talked about dosing about
selection ive talked a little bit about
some of these other issues but
there are important drugs where theres
still additional information needed so
for example the the opiate
pain control medicines all have genetics
that are involved in their metabolism
theres clinical uh studies uh not
randomized trials but uh clinical uh
studies that have looked at these
genetic variants and seen some effect
but there havent been those prospective
randomized trials to really look in and

say

can we use this genetic information
to guide therapy better select therapy
in some way and with the opiate epidemic
thats occurring in many countries
around the world
its certainly an important factor that
if we can more rationally use these
medicines to control pain
and to hopefully manage their their
misuse uh then were going to be in a
much better shape as a society and

individual patients will have better effects

same with the antimedics the these
antimedics are used of course to
prevent nausea and vomiting from cancer
drugs but a more common use is every
patient getting general anesthesia for
surgical procedures for any other type
of procedure

well well receive one of these
antimedics to prevent
nausea and vomiting
again genetic variation is occurring and
can be important for the effect of these

drugs

heres a chance to try to better
understand this
try to use this as ways of guiding

there are

therapy

multiple millions of patients getting surgical procedures every year just with hip replacements its almost 00 000 patients a year in the united states

alone

all of those getting general anesthesia

all those could benefit from genetic
guided antinausea nausea antivomiting
medicines and so a real opportunity that
has not yet been realized to affect the
dosing or dose selection of very
commonly used medicines across many
different therapeutic areas

now

towards the closing here i want to just
mention one thing about how this
information can also be used at a public

health level

when were looking at
major com major influences on health
care modern medical therapy has really

been a key component of improving health

now clean water

other other factors have also been very

important but

access to medicines is an important part

of this

and and choosing these medicines is a major part of most developing countries

the decision making

and so

basing a decision based on access and

costs is important

but if

a if a

drug b is something like familiarity or
clinical consensus thats fine but its
not necessarily the best way of trying
to select medicines and so can we do
more and so for example warfarin is a
medicine that youve heard about in some
of the other presentations in this

series

used across the world
including over million prescriptions
for this medicine in the united states

alone

its a medicine that is metabolized by a
number of different factors number
different p0 enzymes sip enzymes as
shown here also genetic variation in the

target the vitamin k

oxidoreductase

is also a factor and so one can look at these genetic features and then one can put it into a guideline so theres

something called warfarindosingorg

as one example

of a of a tool thats out there where
one can put in features such as age race
weight etc put in genetic information

press the button

and then end up with recommendations on
the dose that should be used per week
to to guide an individual patient
and thats great that information could

be quite useful

but if there is no genetic information available for your country and youre trying to look at differences what can

you do

well one can go in and
one has done the pharmacogenetics for
every nation initiative or pgenie as

its also known

went and looked at large cohorts from
across many different world populations
looked at the genetic information as
well as average body size et cetera was
able to were able to look at what is
the average uh predicted dose based on
these algorithms in a given country and

so what whats shown in green is is uh
european countries uh whats shown in in
um in blue is asian countries uh and
whats shown in in red are african
countries and hopefully you can see from
here that most african countries um have
a average dose of warfarin needed to
give anticlotting activity that is much
higher than seen in the european

populations

most asian countries need a much lower dose compared to the european countries now theres exceptions such as india where there are more caucasoid mixed population so its more close to the europeans also south africa where the population that was studied included a number of the different uh african tribes as well as the um socalled uh mixed or or as its known in south africa colored populations as as well as the afrikaans white populations and so you see this this lower dose but a lot of variation happening across the world um in in even within subcontinents um for which

dose of warfarin one might pick

for an average dose within a country

and then whats shown here

is the algorithms from there were

published in the new england journal

applied to

a chinese and japanese population a us and mexican population and a ghanaian

and

nigerian population showing that uh not what i showed before that the average dose needed for uh in this case two west african populations are much higher than that seen in the us white population whereas the asian in this case japanese and chinese need a much lower dose but you can see within every one of the populations theres a big spread there are people within ghana that need a lower than us average dose and there are people that need extremely high doses there are people within the

us need a

a dose thats lower than needed in asia and there are people that need a dose and these are in the us white

population need a dose thats higher
than seen in africa and so its not as
simple as a given continent a given
color a given selfdeclared
race or ethnicity but really needing
individual patient level data will help

us

guide the therapy in a much more useful
way but in the meantime one can start
with a lower dose in asia and a higher
dose in africa and then individualize as
needed because of these these average

features

so i want to finish off with this this

slide

thats thats really reminding us that
there is still a lot of work to be done
we dont know enough about blood level
guided dosing

if blood level guided dosing can be
applied in much
much beyond just antidepressants
that is a feature that is underutilized
and can be can be
a part of the solution

we also have very few examples where

genetic variation in

transporter proteins

or even in pharmacodynamic examples
are are present and are to the point
where we can use them routinely in
clinical care and so i think thats an
important uh research area where we need
to be doing more work in the context of
transporters and in the the receptors
the targets the pharmacodynamic factors
to really be able to better understand
these areas and better individualize

therapy

options are almost are often limited
based on available dosage forms
so you may be able to determine based on
a body weight an age a genetic test
exactly how many milligrams a patient

needs

but the medicines might be only
available in 00 milligram tablets and
maybe in capsules that cant be broken
in half uh to reduce the dosing and so
we need to be thinking about their
practical dosage forms in terms of how
we individualize therapy were not going

to be able to dial in the exact
milligram of a medicine for an
individual patient every single time
sometimes it might be people need one
tablet or two tablets or one tablet or a

half a tablet

that level of precision still is better
than our current approach
but is certainly not to the point where
wed like to be but the practicalities
of available dosage forms have to be

also most of the examples are with oral medicines

taken into account

and thats great a lot of medicines
nowadays are oral pills tablets
capsules

and other ways of administering

medications that really need to be

studied as well because were not just

choosing the dose but also whether we

pick a different medicine these genetic

factors can be very important in terms

of that and need to be factored in

and then lastly we need more data on the

effect of pharmacogenetically guided dosing

on the adherence consequences
on economic consequences and on outcome
consequences i i showed you some data
for the antidepressants where there was

superior outcome

i showed you

i hinted at some data for the
thyroipurin drugs where there is equal
outcome and reduction in toxicity but
theres very little data on whether
genetic information can cause someone to
take a medicine at the regular

prescribed rate

rather than

forgetting about it or deciding they
just werent going to take it for a
while so can we increase adherence to a
a medication plan based on genetics
can we show for other medicines as we
did for voroconazole that genetic guided
therapy is going to allow you to have

and

better effectiveness

reduce the cost both the cost to the

health system

and the cost out of pocket to an individual patient so we have a lot to be done still to optimize these these examples but hopefully ive been able to

show you

that there are

many examples already where pharmacogenetic information can help us choose the dosage the the selection of medicines prior to ever treating a patient so that we can try to get the right uh the right medicine at the right dose for this patient from the very start we have a long way to go before its perfect but it is uh it is approach we can now take so hopefully youve enjoyed this uh this presentation uh if you have any uh questions on this presentation feel free to contact the moderators uh of this uh of this series and we hope that you tune in for the other uh presentations in this series and please feel free to give comments on how we can improve it and meet your needs as we go forward thank

you very much

you